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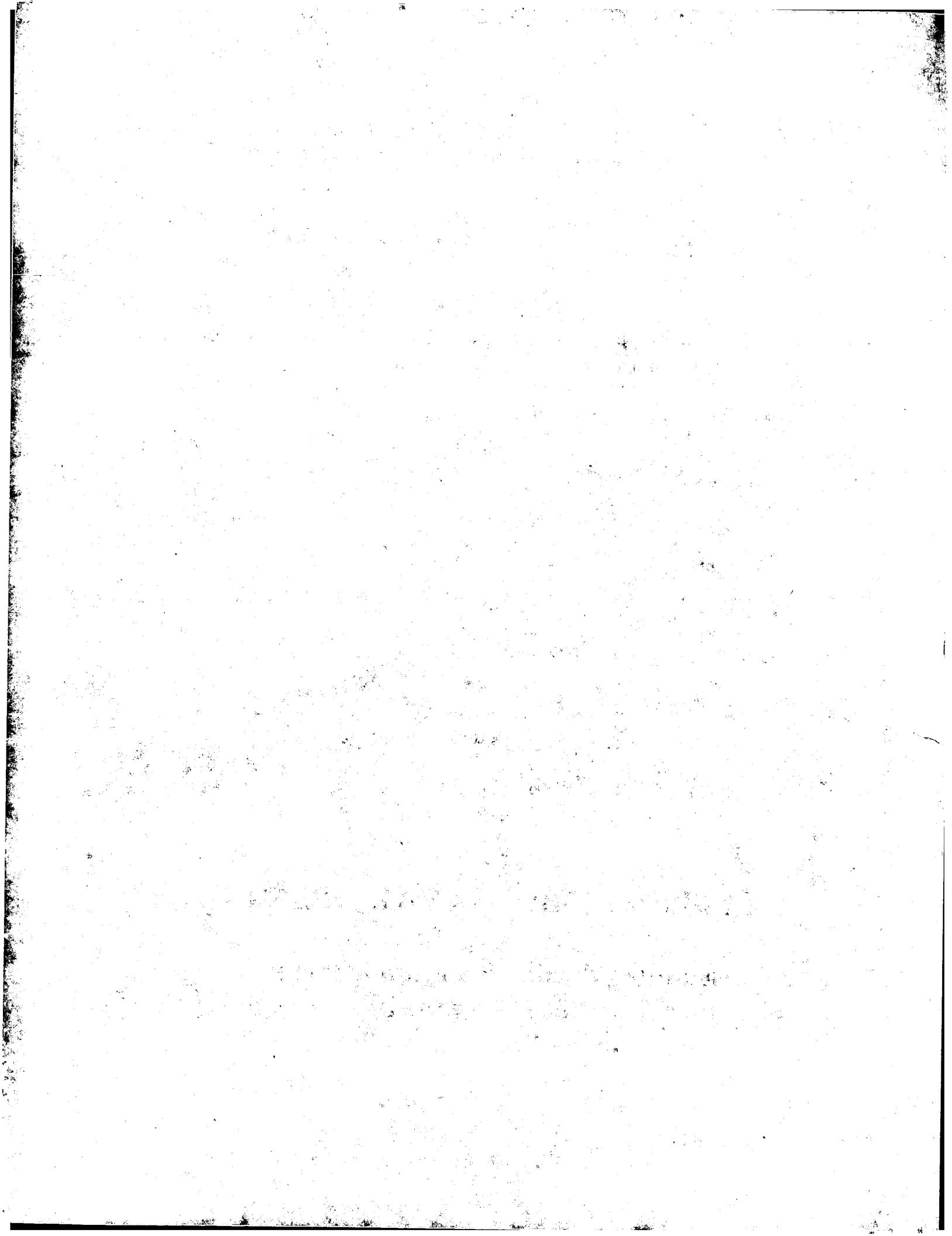
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10/0001433
12/29/03



(19)

Europäisches Patentamt

European Patent Office

Office européen des brevets



(11) EP 0 773 288 A2

(12)

EUROPEAN PATENT APPLICATION

(43) Date of publication:
14.05.1997 Bulletin 1997/20

(51) Int. Cl.⁶: C12N 15/00, A01K 67/027,
C12N 15/90, C12N 5/10,
C07K 16/00, C12N 15/08

(21) Application number: 96113839.3

(22) Date of filing: 29.08.1996

(84) Designated Contracting States:
AT BE CH DE DK ES FI FR GB GR IE IT LI LU MC
NL PT SE

(30) Priority: 29.08.1995 JP 242340/95
15.02.1996 JP 27940/96

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Remarks:

The applicant has subsequently filed a sequence
listing and declared, that it includes no new matter.

(54) Chimeric animal and method for producing the same

(57) Chimeric non-human animals retaining foreign chromosomes or fragments thereof and expressing the genes on the foreign chromosomes or fragments, and their progenies; a method for producing the chimeric non-human animals; tissues and cells derived from the chimeric non-human animals and their progenies; hybridomas prepared by fusion of said cells with myeloma cells; pluripotent cells containing foreign chromosomes or fragments thereof; a method for producing the pluripotent cells; use of the pluripotent cells for producing chimeric non-human animals; and a method for producing a biologically active substance, which comprises expressing the gene on a foreign chromosome or a fragment in the chimeric non-human animal or its progeny, or a tissue or a cell thereof, and recovering the biologically active substance as the expression product.

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Description**TECHNICAL FIELD**

5 The present invention relates to chimeric non-human animals, a method for producing the same and a method for using the same. The present invention allows chimeric non-human animals to retain a foreign giant DNA fragment(s) of at least 1 Mb and to express the gene(s) on such a fragment(s), which was impossible heretofore. Hence, the following becomes possible by using the method.

10 • Production of animals which retain and express a full length of a gene encoding a biologically active substance, for example, a full length of human antibody gene. The biologically active substance, for example, a human-type antibody is useful as a pharmaceutical product.
 • Analysis of functions of human giant genes (e.g., histocompatibility antigen, dystrophin, etc.) in animals.
 • Production of model animals with human dominant hereditary disease and a disease due to chromosomal aberration.

15

BACKGROUND ART

20 Techniques of expressing foreign genes in animals, that is, techniques of producing transgenic animals are used not only for obtaining information on the gene's functions in living bodies but also for identifying DNA sequences that regulate the expression of the genes (e.g., Magram et al., *Nature*, 315:338, 1985), for developing model animals with human diseases (Yamamura et al., "Manual of model mice with diseases" published by Nakayama Shoten, 1994), for breeding farm animals (e.g., Muller et al., *Experientia*, 47:923, 1991) and for producing useful substances with these animals (e.g., Velander et al., *P.N.A.S.*, 89:12003, 1992). Mice have been used the most frequently as hosts for gene transfer. Since mice have been studied in detail as experimental animals and the embryo manipulating techniques for mice have been established, they are the most appropriate kind of mammals for gene transfer.

25 Two methods are known for transferring foreign genes into mice. One is by injecting DNA into a pronucleus of a fertilized egg (Gordon et al., *P.N.A.S.*, 77:7380, 1980). The other is by transferring DNA into a pluripotent embryonic stem cell (hereinafter referred to as "ES cell") to produce a chimeric mouse (Takahashi et al., *Development*, 102:259, 1988).
 30 In the latter method, the transferred gene is retained only in ES cell-contributing cells and tissues of chimeric mice whereas it is retained in all cells and tissues of progenies obtained via ES cell-derived germ cells. These techniques have been used to produce a large number of transgenic mice up to now.

35 However, there had been a limit of the size of DNA capable of being transferred and this restricts the application range of these techniques. The limit depends on the size of DNA which can be cloned. One of the largest DNA fragments which have ever been transferred is a DNA fragment of about 670 kb cloned into a yeast artificial chromosome (YAC) (Jakobovits et al., *Nature*, 362:255, 1993). This experiment was carried out by fusing a YAC-retaining yeast cell with a mouse ES cell. Although it is believed that foreign DNA of up to about 2 Mb can be cloned on YAC (Den Dunnen et al., *Hum. Mol. Genet.*, 1:19, 1992), the recombination between homologous DNA sequences occurs frequently in budding yeast cells and therefore, in some cases, a human DNA fragment containing a large number of repeated sequences is difficult to retain in a complete form. In fact, certain recombinations occur in 20-40% of the clones of YAC libraries containing human genomic DNA (Green et al., *Genomics*, 11:584, 1991).

40 In another method that was attempted, a metaphase chromosome from a cultured human cell was dissected under observation with a microscope and the fragment (presumably having a length of at least 10 Mb) was injected into a mouse fertilized egg (Richa et al., *Science*, 245:175, 1989). In the resulting mice, a human specific DNA sequence (Alu sequence) was detected but the expression of human gene was not confirmed. In addition, the procedure used in this method to prepare chromosomes causes unavoidable fragmentation of DNA into small fragments due to the use of acetic acid and methanol in fixing the chromosome on slide glass and the possibility that the injected DNA exists as an intact sequence is small.

45 In any event, no case has been reported to date that demonstrates successful transfer and expression in mice of uninterrupted foreign DNA fragments having a length of at least 1 Mb.

50 Useful and interesting human genes which are desirably transferred into mice, such as genes for antibody (Cook et al., *Nature Genetics*, 7: 162, 1994), for T cell receptor (Hood et al., *Cold Spring Harbor Symposia on Quantitative Biology*, Vol. LVIII, 339, 1993), for histocompatibility antigen (Carrol et al., *P.N.A.S.*, 84:8535, 1987), for dystrophin (Den Dunnen et al., *supra*) are known to be such that their coding regions have sizes of at least 1 Mb. Since human-type antibodies are important as pharmaceutical products, the production of mice which retain and express full lengths of genes for human immunoglobulin heavy chains (~1.5 Mb, Cook et al., *supra*), and light chain κ (~3 Mb, Zachau, *Gene*, 135:167, 1993), and light chain λ (~1.5 Mb, Frippiat et al., *Hum. Mol. Genet.*, 4:983, 1995) is desired but this is impossible to achieve by the state-of-the-art technology (Nikkei Biotec, July, 5, 1993).

55 Many of the causative genes for human dominant hereditary disease and chromosomal aberration which causes

congenital deformity (Down's syndrome, etc.) have not been cloned and only the information on the approximate location of the genes on chromosome is available. For example, when a gene of interest is found to be located on a specific G band, which is made visible by subjecting a metaphase chromosome to Giemsa staining, the G band has usually a size of at least several Mb to 10 Mb. In order to transfer these abnormal phenotypes into mice, it is necessary to transfer chromosomal fragments of at least several Mb that surround the causative genes, but this is also impossible with the presently available techniques.

Hence, it is desired to develop a technique by which a foreign DNA longer than the heretofore critical 1 Mb can be transferred into a mouse and expressed in it.

DNA longer than 1 Mb can be transferred into cultured animal cells by the techniques available today. Such transfer is carried out predominantly by using a chromosome as a mediator. In the case of human, chromosomes have sizes of about 50-300 Mb. Some methods for chromosome transfer into cells have been reported (e.g., McBride et al., P.N.A.S., 70:1258, 1973). Among them, microcell fusion (Koi et al., Jpn. J. Cancer Res., 80:413, 1989) is the best method for selective transfer of a desired chromosome. The microcell is a structural body in which one to several chromosomes are encapsulated with a nuclear membrane and a plasma membrane. A few chromosomes (in many cases, one chromosome) can be transferred by inducing a microcell with an agent that inhibits the formation of spindle in a specific kind of cell, separating the microcell and fusing it with a recipient cell. The resulting libraries of monochromosomal hybrid cells containing only one human chromosome have been used for mapping known genes and specifying the chromosomes on which unknown tumor-suppressor genes and cellular senescence genes exist (e.g., Saxon et al., EMBO J., 5:3461, 1986). In addition, it is possible to fragment a chromosome by irradiating a microcell with γ -rays and to transfer part of the fragments (Koi et al., Science, 260:361, 1993). As described above, microcell fusion is considered to be an appropriate method for transferring DNA of at least 1 Mb into a cultured animal cell.

The expectation that a mouse could be generated from a cultured cell turned to a real fact when the ES cell which has stable pluripotency was discovered (Evans et al., Nature, 292:154, 1981). Foreign genes, various mutations and mutations by targeted gene recombination could be introduced into the ES cell, making it possible to perform a wide variety of genetic modifications in mice (e.g., Mansour et al., Nature, 336:348, 1988). As long as the transfer of giant DNA is concerned, it has been believed that the size of the aforementioned foreign DNA fragment which can be cloned into a YAC vector is the upper limit. The prior art technology of chromosome transfer for introducing a longer DNA into cultured cells has never been applied to gene transfer into mice and this has been believed to be difficult to accomplish (Muramatsu et al., "Transgenic Biology", published by Kodansha Scientific, p.143-, 1989).

The reasons are as follows.

- The transfer of a human chromosome into a mouse ES cell of a normal karyotype as a recipient cell would be a kind of transfer of chromosomal aberration. Up to now, it has been believed that genetic aberration at chromosomal levels which is large enough to be recognizable with microscopes is generally fatal to the embryogeny in mice (Gropp et al., J. Exp. Zool., 228:253, 1983 and Shinichi Aizawa, "Biotechnology Manual Series 8, Gene Targeting", published by Yodobashi, 1995).
- Available human chromosomes are usually derived from finitely proliferative normal fibroblasts or differentiated somatic cells such as cancer cells and the like. It was believed that if a chromosome derived from such a somatic cell was transferred into an undifferentiated ES cell, the transferred chromosome might cause differentiation of the ES cell or its senescence (Muller et al., Nature, 311:438, 1984; Sugawara, Science, 247:707, 1990).
- Only few studies have been reported as to whether a somatic cell-derived chromosome introduced into an early embryo can function in the process of embryonic development as normally as a germ cell-derived chromosome to ensure the expression of a specific gene in various kinds of tissues and cells. One of the big differences between the two chromosomes is assumed to concern methylation of the chromosomal DNA. The methylation is changed according to differentiation of cells and its important role in the expression of tissue-specific genes has been suggested (Ceder, Cell, 53:3, 1988). For example, it has been reported that if a methylated DNA substrate is introduced into a B cell, the methylated DNA is maintained after replication and suppresses a site-directed recombination reaction which is essential to the activation of an antibody gene (Hsieh et al., EMBO J., 11:315, 1992). In addition, it was reported that higher levels of de novo methylation occurred in established cell lines than in vivo (Antequera et al., Cell, 62:503, 1990). On the basis of the studies reported, it could not be easily expected that an antibody gene in a human fibroblast or a human-mouse hybrid cell which was likely to be methylated at a high level would be normally expressed in a mouse B cell.

It should be noted that there are two related reports of Illmensee et al. (P.N.A.S., 75:1914, 1978; P.N.A.S., 76:879, 1979). One report is about the production of chimeric mice from fused cells obtained by fusing a human sarcoma cell with a mouse EC cell and the other is about the production of chimeric mice from fused cells obtained by fusing a rat liver cancer cell with a mouse EC cell. Many questions about the results of the experiments in these two reports were pointed out and thus these reports are considered unreliable (Noguchi et al., "Mouse Teratoma", published by Rikogakusha, Section 5, 1987). Although it has been desired to perform a follow-up as early as possible, as of today

when 17 years have passed since the publication of these reports, successful reproduction of these experiments has not been reported. Hence, it is believed that foreign chromosomes cannot be retained and the genes on the chromosomes cannot be expressed in mice by the method described in these reports.

Under these circumstances, it has been believed that it is difficult to transfer a giant DNA such as a chromosomal fragment and express it in an animal such as mouse. Actually, no study has been made about this problem since the filimensee's reports.

Therefore, an object of the present invention is to provide chimeric non-human animals which retain foreign chromosomes or fragments thereof and express genes on the chromosomes or fragments, and their progenies, and a method for producing the same.

It is also an object of the present invention to provide pluripotent cells containing foreign chromosomes or fragments thereof and a method for producing the pluripotent cells.

Another object of the present invention is to provide tissues and cells derived from the chimeric non-human animals and their progenies.

A further object of the present invention is to provide hybridomas prepared by fusing the cells derived from the chimeric non-human animals and their progenies with myeloma cells.

A still further object of the present invention is to provide a method for producing a biologically active substance that is an expression product of the gene on a foreign chromosome or a fragment thereof by using the chimeric non-human animals or their progenies, or their tissues or cells.

20 DISCLOSURE OF THE INVENTION

As a result of the various studies conducted to achieve the above objects, the inventors succeeded in transferring chromosomes or fragments thereof derived from human normal fibroblast cells into mouse ES cells and obtaining clones which were capable of stable retention of the chromosomes or fragments. Moreover, they produced from these ES clones those chimeric mice which retained human chromosomes in normal tissues and which expressed several human genes including human antibody heavy-chain genes. It has become possible to make that animals retain and express giant DNA fragments by the series of these techniques, although this has been impossible by conventional techniques. The present invention provides a method for producing a chimeric non-human animal, which comprises preparing a microcell containing a foreign chromosome(s) or a fragment(s) thereof and transferring the foreign chromosome(s) or fragment(s) into a pluripotent cell by fusion with the microcell. The present invention also provides chimeric non-human animals and their progenies that retain foreign chromosomes or fragments thereof and which express the genes on the foreign chromosomes or fragments. In addition, the present invention provides tissues and cells derived from the chimeric non-human animals and progenies. The present invention also provides hybridomas prepared by fusion of the cells with myeloma cells. Furthermore, the present invention provides a method for producing a pluripotent cell containing a foreign chromosome(s) or a fragment(s) thereof, which comprises preparing a microcell containing a foreign chromosome(s) or a fragment(s) thereof and transferring the foreign chromosome(s) or fragment(s) into a pluripotent cell by fusion with the microcell. The present invention also provides pluripotent cells that contain foreign chromosomes or fragments thereof and use thereof for producing chimeric non-human animals. Moreover, the present invention provides a method for producing a biologically active substance, which comprises expressing the gene on a foreign chromosome or a fragment in the chimeric non-human animal or its progeny, or a tissue or a cell thereof.

BRIEF DESCRIPTION OF THE DRAWINGS

Fig. 1 shows the results of PCR analysis of an A9 cell retaining human chromosome #2 (fragment).

Fig. 2 shows that human chromosome #22 (fragment) is retained in an E14 drug resistant cell (PCR analysis).

Fig. 3 is a photograph of electrophoresis patterns showing that human L1 sequence is retained in a chimeric mouse produced from a human chromosome #22-transferred ES cell (Southern analysis).

Fig. 4 is a photograph of electrophoresis patterns showing the presence of a human chromosome in organs of a human chromosome #22 transferred chimeric mouse (PCR analysis).

Fig. 5 is a photograph of electrophoresis patterns showing the results of the expression of human genes in a human chromosome #22 transferred chimeric mouse (RT-PCR).

Fig. 6 is a photograph of electrophoresis patterns showing the results of the expression of human genes in organs of a human chromosome #22 transferred chimeric mouse (RT-PCR).

Fig. 7 shows that human chromosome #4 (fragment) is retained in an E14 drug resistant cell (PCR analysis).

Fig. 8 is a photograph of electrophoresis patterns showing the detection of human L1 sequence in a human chromosome #4-transferred E14 cell clone (Southern analysis).

Fig. 9 is a photograph of electrophoresis patterns showing that human L1 sequence is retained in a chimeric mouse produced from a human chromosome #4-transferred ES cell (Southern analysis).

Fig. 10 shows that human chromosome #14 (fragment) is retained in a TT2 drug resistant cell (PCR analysis).

Fig. 11 is a photograph of electrophoresis patterns showing the presence of a human chromosome in organs of a chimeric mouse produced from a human chromosome #14 transferred ES cell (PCR analysis).

Fig. 12 shows the results of a test on a tail-derived fibroblast cell for resistance to G418.

Fig. 13 shows the concentration of human antibody IgM in a serum of a human serum albumin (hereinafter referred to as "HSA")-immunized chimeric mouse (ELISA).

Fig. 14 shows the concentration of human antibody IgG in a serum of an HSA-immunized chimeric mouse (ELISA).

Fig. 15 shows the results of ELISA of hybridoma clone H4B7 capable of producing human IgM.

Fig. 16 is a photograph of the results of FISH analysis of a mouse ES cell clone (TT2 cell clone PG15) retaining partial fragments of human chromosomes #2 and 14.

Fig. 17 shows that the antibody titer of anti-HSA human IgG is increased in a serum of an HSA-immunized chimeric mouse.

Fig. 18 shows that the antibody titer of anti-HSA human Ig κ is increased in a serum of an HSA-immunized chimeric mouse.

Fig. 19 is a photograph of electrophoresis patterns showing the detection of human L1 sequence in a human chromosome #22-transferred TT2 cell clone (Southern analysis).

Fig. 20 shows that the antibody titer of anti-HSA human Ig λ is increased in a serum of an HSA-immunized chimeric mouse.

Fig. 21 shows that a partial fragment of human chromosome #2 is retained in a progeny of a chimeric mouse into which a partial fragment of a human chromosome #2 was transferred (PCR analysis).

Fig. 22 shows the presence of a cell expressing human μ chain on the cell surface in a spleen of a human chromosome #14-transferred chimeric mouse (flow cytometry analysis).

DETAILED DESCRIPTION

The present invention will now be described in detail.

A non-human animal that retains a human chromosome(s) or a fragment(s) thereof and which expresses the gene on the chromosome(s) or fragment(s) thereof can be produced by

(1) preparing a chromosome donor cell which retains a labeled human chromosome or a fragment thereof;

(2) transferring the human chromosome or fragment thereof into a nonhuman animal pluripotent cell by microcell fusion;

(3) producing a chimeric non-human animal from the cell; and

(4) confirming that the human chromosome is retained in the chimeric non-human animal and that a human gene is expressed.

In this procedure, a mouse is used as a non-human animal that retains a human chromosome or a fragment thereof and which expresses the gene on the chromosome or fragment thereof (the mouse is hereinafter referred to as a "human chromosome transferred mouse").

The term "human chromosome" means a naturally occurring complex which consists of nucleic acids and proteins that are derived from human cells. There are 46 normal human chromosomes of 23 kinds (24 kinds in male), each of which contains DNAs of about 50-300 Mb in size. In the present invention, the human chromosome includes not only partial fragments which can be stably replicated and segregated as independent chromosomes but also fragments that are translocated on mouse chromosomes and which are retained stably. The size of the DNA is usually at least 1 Mb and in some cases, it is smaller than 1 Mb. The feature of the present invention resides in that a mouse can retain and express the foreign gene on a foreign chromosome as a mediator without treatments such as cloning in an *E. coli* or yeast cell, or extraction of the DNA from a cell.

The term "human chromosome transferred mouse" means a mouse retaining a human chromosome(s) or a fragment(s) thereof in all or part of its normal somatic cells. The mouse expresses the gene(s) on a human chromosome(s) or a fragment(s) thereof in all or part of its normal somatic cells.

(1) Preparation of a chromosome donor cell which retains a labeled human chromosome or a fragment thereof

A desired chromosome donor cell 1) retains a human chromosome(s) labeled with a marker also available for selection of recipient cells; 2) does not contain other human chromosomes; and 3) has a higher ability to form a microcell.

Any human-derived cell lines, cancer cells and primary culture cells can be used as materials for providing human chromosomes. Among them, normal fibroblast cells are suitable because they have a low possibility of abnormality such as deletion and amplification of chromosomes and can be readily cultured.

As for 1), human cells can be transformed with vectors that express genes for markers such as drug-resistance

(e.g., G418-, puromycin-, hygromycin- or blasticidin-resistance). Promoters operating efficiently not only in human cells but also in recipient cells such as mouse ES cells are desirably used to regulate the expression of the marker used. For this purpose, herpes simplex virus thymidine kinase promoter linked with SV 40 enhancer (Katoh et al., Cell Struct. Funct., 12:575, 1987), mouse PGK-1 promoter (Soriano et al., Cell, 64:693, 1991) and the like can be used. A library of human cell transformants in which the introduced marker genes have been inserted into 46 human chromosomes of 23 kinds at random can be prepared by transformation through electroporation (Ishida et al., "Cell Technology Experiment Manual", published by Kodansha, 1992) and the like and subsequent selection of transformants.

As for 3), since many human normal cells have a very low ability to form microcells, the whole cell of the transformant may be fused with a cell having a high ability to form microcells such as mouse A9 cell (Oshimura, M., Environ. Health Perspect., 93:57, 1991) so as to provide the transformed cell with an ability to form microcells. It is known that in mouse-human hybrid cells, human chromosomes selectively disappear. The fused cell selected by the marker can retain stably the marked human chromosome.

In order to meet the condition of 2), it is desired to obtain a microcell from the fused cell and fuse it again with a mouse A9 cell. In this case, too, most of the cells selected by the marker will meet the three conditions 1), 2) and 3) above. The marked human chromosomes can be identified in the finally obtained mouse-human monochromosomal hybrid cells by PCR (Polymerase Chain Reaction, Saiki et al., Science, 239:487, 1988), Southern blot analysis (Ausubel et al., Current protocols in molecular biology, John Wiley & Sons, Inc., 1994), FISH analysis (Fluorescence In Situ Hybridization, Lawrence et al., Cell, 52: 51, 1988) and the like. If the transfer of a specified chromosome is desired, the above procedures are applied to each of many human cell transformant clones to select a clone in which a chromosome of interest is marked. Alternatively, the above procedures are applied to a mixture of human cell transformant clones and the identification of human chromosomes is carried out on a large number of the resulting mouse-human monochromosome hybrid cells.

In addition, a marker gene can be inserted into a desired site by homologous recombination of a specific DNA sequence on the chromosome which is to be transferred (Thomas et al., Cell, 51:503, 1987).

A microcell prepared from the mouse-human hybrid cell may be irradiated with γ -rays such that the marked human chromosome is fragmented and transferred into a mouse A9 cell. Even if the microcell is not irradiated with γ -rays, a partially fragmented human chromosome may be transferred at a certain frequency. In these cases, the resulting microcell fused clones retain partial fragments of the marked human chromosomes. These clones can be used when it is desired to transfer the partial fragments into recipient cells.

30 (2) Transfer of the human chromosome or fragment thereof into a mouse pluripotent cell

It has been reported to date that an embryonic carcinoma cell (EC cell, Hanaoka et al., Differentiation, 48:83, 1991), an embryonic stem cell (ES cell, Evans, Nature, 292:154, 1981) or an embryonic germ cell (EG cell, Matsui et al., Cell, 70:841, 1992) that are derived from various strains of mice contribute to the normal somatic cells in mice, or are capable of the production of chimeric mice, by injection into or coculturing with a mouse early embryo. ES and EG cells have a very high ability in this respect and in many cases, they also contribute to germ cells thereby making it possible to produce progenies derived from the cells. EC cells can be obtained predominantly from teratocarcinoma; ES cells from the inner cell masses of blastocysts; and EG cells from primordial germ cells appearing at the early stage of embryogeny. These cell lines and their mutants, and any undifferentiated cells that are capable of differentiation into all or part of the normal somatic cells in mice can be used as recipient cells for the transfer of human chromosome in the present invention.

The microcells prepared from the human chromosome donor cells or the microcells irradiated with γ -rays can be used as materials for the transfer of human chromosomes into the recipient cells. The human chromosome can be transferred into the recipient cell through fusion of the recipient cell with the microcell by the method described in Motoyuki Shimizu, "Cell Technology Handbook", published by Yodobashi, 1992. The microcell donor cells retain markers by which human chromosomes or fragments thereof can be selected in the recipient cells. The clone containing a gene, a chromosome or a fragment of interest can be selected by PCR, Southern blot analysis, FISH method or the like in the same manner as in (1), thus all kinds of human chromosomes or fragments thereof can be transferred. Moreover, if several chromosomes or fragments thereof which contain different selection markers are transferred sequentially, a recipient cell retaining these chromosomes or fragments at the same time can be obtained. In addition, clones having an increased number of the transferred chromosome can be selected from the clones into which the human chromosome has been transferred. Such selection can be accomplished by increasing the concentration of a selection drug to be added to a culture medium.

55 In order to determine whether the recipient cell selected by the marker (e.g., G418 resistance) on the human chromosome retains the whole or part of the chromosome retained by the donor cell, the following confirmative techniques may be employed: South blot analysis using the genomic DNA extracted from the selected recipient cell, with a human specific repeated sequence (L1, Alu, etc.: Korenberg et al., Cell, 53:391, 1988) or a human gene used as a probe; and chromosome analysis such as PCR method using a human gene specific primer or FISH method using a

human chromosome specific probe (Lichter et al., Human Genetics, 80:224, 1988).

(3) Production of a chimeric mouse from the human chromosome transferred ES cell

5 The method described in Shinichi Aizawa, "Biotechnology Manual Series 8, Gene Targeting", published by Yodo-sha, 1995 may be used to produce chimeric mice from the ES cell clone obtained in (2). In selecting factors for efficient production of chimeric mice, such as the developmental stage of the host embryo and its strain, it is desired to employ the conditions already reviewed for the respective ES cell clones. For example, 8 cell stage embryos derived from Balb/c (albino, CREA JAPAN, INC.) or ICR (albino, CREA JAPAN, INC.) are desirably used for CBAXC57BL/6 F1-derived TT2 cell (agouti, Yagi et al., Analytical Biochemistry, 214:70, 1993).

10 (4) Confirmation of the retention of the human chromosome in the chimeric mice and the expression of a human gene

15 The contribution of the ES cells in mice produced from the embryos into which ES cells were injected can be roughly judged by the color of their coat. However, it should be noted that the total absence of contribution to the coat color does not always lead to the conclusion that there is no contribution to other tissues. The detailed information on the retention of the human chromosome in various tissues of the chimeric mice can be obtained by Southern blot analysis using the genomic DNA extracted from various tissues, by PCR or the like.

20 The expression of the gene on the transferred human chromosome can be confirmed by the following methods.

25 The expression of mRNA transcribed from the human chromosome can be detected by RT-PCR method or northern blotting (Ausubel et al., supra) using RNAs derived from various tissues (Kawasaki et al., P.N.A.S., 85:5698, 1988). The expression at the protein level can be detected by enzyme immunoassay using an anti-human protein antibody that is rendered minimal in its ability to enter into a cross reaction with mouse homologous proteins (ELISA, Toyama and Ando, "Monoclonal Antibody Experiment Manual", published by Kodansha Scientific, 1987; Ishikawa, "Enzyme immunoassay with Superhigh Sensitivity", published by Gakkai Shuppan Center, 1993), western blotting (Ausubel et al., supra), isozyme analysis utilizing the difference in electrophoretic mobility (Koi et al., Jpn. J. Cancer Res., 80:413, 1989) or the like. The retention of the human chromosome in the chimeric mice and the expression of the gene on the human chromosome can be confirmed by the appearance of the cells expressing a drug resistance marker gene in primary culture cells derived from the chimeric mice.

30 For example, human IgM, IgG, IgA and the like in sera of the chimeric mice which are produced from ES cells retaining human chromosome #14 on which a gene for human immunoglobulin heavy chain exists can be detected by enzyme immunoassay using an anti-human Ig antibody that is rendered minimal in its ability to enter into cross reaction with mouse antibody. Hybridomas capable of producing a human immunoglobulin heavy chain can be obtained by ELISA screening of hybridomas prepared by immunizing the chimeric mouse with a human-derived antigen (e.g., HSA) and fusing the spleen cells of the immunized mice with mouse myeloma cells (Toyama and Ando, "Monoclonal Antibody Experiment Manual", published by Kodansha Scientific, 1987).

35 The method for producing a chimeric non-human animal of the present invention has been explained above with reference to the case of a mouse retaining a human chromosome(s) or a fragment(s) thereof and expressing the gene(s) on the chromosome(s) or fragment(s). In the present invention, chromosomes or fragments thereof to be transferred into chimeric non-human animals are not limited to those derived from humans but include any foreign chromosomes and fragments thereof. The foreign chromosome and fragment thereof can be transferred into pluripotent cells and the gene on the chromosome and fragment thereof can be expressed in the chimeric non-human animal. The origin of the foreign chromosome and fragment thereof is not particularly limited. Other kinds of chimeric animals such as chimeric rat and pig can be produced by the method of the present invention. ES cells or ES-like cells derived from animals other than mouse were established with rat (Iannaccone et al., Dev. Biol., 163, 288, 1994), pig (Wheeler et al., Reprod. Fertil. Dev., 6, 563, 1994) and bovine (Sims et al., Proc. Natl. Acad. Sci. USA, 91, 6143-6147, 1994) and attempts have been made on cyprinodont, chicken and the like ("Transgenic Animal", Protein • Nucleic Acid • Enzyme, October, 1995, Special Issue, published by Kyoritsu Shuppan). It is known that sheep is developed normally from an unfertilized egg transplanted with the nucleus from ES-like cell (ED cell) or epithelial-like cell obtained by subcultivation of the ES-like cell through at least 10 generations (Campbell et al., Nature, 380, 64, 1996). These ES cells and ES-like cells can be used as recipient cells in the transfer of foreign chromosomes to produce chimeric non-human animals retaining the foreign chromosomes or fragments thereof and expressing the genes on the chromosomes or fragments thereof in the same manner as in the case of mouse.

40 50 55 If an ES cell retaining a foreign chromosome(s) or a fragment(s) thereof is differentiated to a germ cell in the chimeric non-human animal, reproduced progenies will retain the transferred chromosome(s) or fragment(s) thereof and express the gene(s) on the chromosome(s) or fragment(s) thereof.

The chimeric non-human animals or their progenies can be used to express the gene on the foreign chromosome or fragment thereof and to recover the expression product, thereby producing a biologically active substance. More specifically, the chimeric non-human animals or their progenies can be bred under the conditions for expressing the gene

on the foreign chromosome or fragment thereof to recover the expression product from the blood, ascites and the like of the animals. Alternatively, the tissues or cells of the chimeric non-human animal, or immortalized cells derived therefrom (e.g., hybridomas immortalized by fusion with myeloma cells) can be cultured under the conditions for expressing the gene on the foreign chromosome or fragment thereof and the expression product is thereafter recovered from the culture. The expression product can be collected by known techniques such as centrifugation and purified by known techniques such as ammonium sulfate fractionation, partition chromatography, gel filtration chromatography, adsorption chromatography, preparative thin-layer chromatography and the like. The biologically active substance includes any kinds of substances encoded on foreign chromosomes, for example, antibodies, particularly human antibodies.

The chimeric mice or their progenies that retain human chromosomes #2, 14 and/or 22 (or fragments thereof) which can be produced by the method of the present invention can retain the greater part of the functional sequences of respective genes for human antibody heavy chain on chromosome #14, light chain on chromosome #2 and light chain λ on chromosome #22. Hence, they can produce a wide repertory of antibodies which are more similar to human antibody repertory, compared with known transgenic mice into which parts of human antibody gene have been transferred by using yeast artificial chromosomes and the like (Green et al., Nature Genetics, 7, 13, 1994; Lonberg et al., Nature, 368, 856, 1994). The chimeric mice and their progenies retaining chromosomes #2+#14 or #22+#14 or fragments thereof can produce complete human antibodies both heavy and light chains of which are derived from humans. These mice can recognize human-derived antigens as foreign substances to cause an immunoreaction with the antigens, thereby producing antigen-specific human antibodies. These properties can be utilized to produce human monoclonal and polyclonal antibodies for therapeutic treatments (Green et al, supra; Longberg et al., supra)

The present invention will now be explained in greater detail with reference to the following examples, which do not limit the scope of the present invention.

Example 1

25 Production of chromosome donor cell retaining human chromosome (fragment) labeled with G418 resistance

Plasmid pSTneoB containing a G418 resistance gene (Katoh et al., Cell Struct. Funct., 12:575, 1987; Japanese Collection of Research Biologicals (JCRB), Deposit Number: VE 039) was linearized with restriction enzyme Sall (TAKARA SHUZO CO., LTD.) and introduced into human normal fibroblast cell HFL-1 (obtained from RIKEN Cell Bank, RCB0251). The HFL-1 cells were treated with trypsin and suspended in Dulbecco's phosphate-buffered saline (PBS) at a concentration of 5×10^6 cells/ml, followed by electroporation using a Gene Pulser (Bio-Rad Laboratories, Inc.) in the presence of 10 μ g of DNA (Ishida et al., "Cell Technology Experiment Procedure Manual", published by Kodansha, 1992). A voltage of 1000 V was applied at a capacitance of 25 μ F with an Electroporation Cell of 4 mm in length (165-2088, Bio-Rad Laboratories, Inc.) at room temperature. The electroporated cells were inoculated into an Eagle's F12 medium (hereinafter referred to as "F12") supplemented with 15% fetal bovine serum (FBS) in 3-6 tissue culture plastic plates (Corning) of 100 mm \varnothing . After one day, the medium was replaced with a F12 supplemented with 15% FBS and containing 200 μ g/ml of G418 (GENENTICIN, Sigma). The colonies formed after 2-3 weeks were collected in 52 groups each consisting of about 100 colonies. The colonies of each group were inoculated again into a plate of 100 mm \varnothing and cultured.

40 Mouse A9 cells (Oshima, Environ. Health Perspect., 93:57, 1991; JCRB 0211) were cultured in Dulbecco's modified Eagle's medium (hereinafter referred to as "DMEM") supplemented with 10% FBS in plates of 100 mm \varnothing . The G418 resistant HFL-1 cells of 52 groups were cultured in F12 supplemented with 15% FBS and 200 μ g/ml of G418 in plates of 100 mm \varnothing . The mouse A9 cells and HFL-1 cells were treated with trypsin and one fourth to one half of both cells were mixed. The mixed cells were inoculated into a plate of 100 mm \varnothing and cultured in a mixture of equal amounts of DMEM containing 10% FBS and F12 containing 15% FBS for a period ranging from a half day to one day. Cell fusion was carried out in accordance with the method described in Shimizu et al., "Cell Technology Handbook", published by Yodosha, p.127, 1992. The cell surface was washed twice with DMEM and then treated sequentially with 2 ml of a PEG (1:1.4) solution for 1 minute and with 2 ml of PEG (1:3) for 1 minute. After the PEG solution was sucked up, and the cells were washed three times with a serum-free DMEM, followed by cultivation in DMEM supplemented with 10% FBS for 1 day. The cells were dispersed by treatment with trypsin and suspended in a double selective medium (10% FBS supplemented DMEM) containing ouabain (1×10^{-5} M, Sigma) and G418 (800 μ g/ml), followed by inoculation in 3 plates of 100 mm \varnothing . After about 3 weeks cultivation, the colonies formed were treated with trypsin to disperse the cells, which were cultured in a selective medium (10% FBS supplemented DMEM) containing G418 (800 μ g/ml).

45 The cells were dispersed by treatment with trypsin and two groups of the cells were collected, followed by cultivation in 6 centrifuge flasks (Coaster, 3025) of 25 cm 2 until the cell density reached 70-80% confluence. The medium was replaced with a medium (20% FBS supplemented DMEM) containing Colcemid (0.05 μ g/ml, Demecolcine, Wako Pure Chemicals Co., Ltd) and the cells were cultured for 2 days to form microcells. After the culture medium was removed, a cytochalasin B (10 μ g/ml, Sigma) solution preliminarily warmed at 37°C was filled in the 25 cm 2 centrifuge flask, which were inserted into an acryl centrifuge container, followed by centrifugation at 34°C at 8,000 rpm for 1 hour. The micro-

cells were suspended in a serum-free medium and purified by passage through a filter. To the mouse A9 cells cultured to 80% confluence in the flask of 25 cm², the purified microcells were added and the two kinds of cells were fused with a PEG solution. The fused cells were cultured in a G418 containing selective medium and colonies formed were isolated. Human chromosomes #2, 4, 14 and 22 retained in the respective clones were identified by the methods described in (1)-(3) below. All other experimental conditions such as operating procedures and reagents were in accordance with Shimizu et al., "Cell Technology Handbook", published by Yodosha, p127-.

(1) PCR analysis

10 The isolated cells were cultured and genomic DNA was extracted from the cells with a Puregene DNA Isolation kit (Gentra System Co.). PCR was performed using the genomic DNA as a template with human chromosome specific primers to select the clones retaining human chromosome #2, 4, 14 or 22. The PCR amplification was conducted with about 0.1 µg of the genomic DNA as a template, using a thermal cycler (GeneAmp 9600, Perkin-Elmer Corp.) in accordance with the method described in Innis et al., "PCR Experiment Manual", published by HBJ Publication Office, 1991.

15 Taq polymerase was purchased from Perkin-Elmer Corp. and the reaction was performed in a cycle of 94°C, 5 minutes and 35 cycles of denaturing at 94 °C , 15 seconds, annealing at 54-57°C , 15 seconds (variable with the primers) and extension at 72°C , 20 seconds. The gene on each chromosome (O'Brien, Genetic Maps, 6th edition, Book 5, Cold Spring Harbor Laboratory Press, 1993) and polymorphic markers (Polymorphic STS Primer Pair, BIOS Laboratories, Inc.; Weissenbach et al., Nature 359: 794, 1992; Walter et al., Nature Genetics, 7:22, 1994) were used as primers. The 20 primers for the genes were prepared on the basis of nucleotide sequences obtained from data bases such as GenBank, EMBL and the like. The names of the polymorphic primers and the sequences of the primers for the genes will be shown for the respective chromosomes in the following examples (#2, Example 1; #4, Example 6, #14, Example 9; #22, Example 2). The following genetic markers and polymorphic makers (Polymorphic STS Primer Pairs: D2S207, D2S177, D2S156 and D2S159, BIOS Laboratories, Inc.) were used to identify chromosome #2.

25 Cκ (immunoglobulin kappa constant) : 5'-TGGAAGGTGGATAACGCCCT (SEQ ID NO:1), 5'-TCATTCTCCTCCAA-CATTAGCA (SEQ ID NO:2)

FABP1 (fatty acid binding protein-1 liver) : 5'-GCAATCGGTCTGCCGGAAAGA (SEQ ID NO:3), 5'-TTGGATCACTT-GGACCCAG (SEQ ID NO:4)

30 Vk3-2 (immunoglobulin kappa variable) : 5'-CTCTCCTGCAGGGCCAGTCA (SEQ ID NO:5), 5'-TGCTGATGGTGA-GAGTGAACTC (SEQ ID NO:6)

Vk1-2 (immunoglobulin kappa variable) : 5'-AGTCAGGGCATTAGCAGTGC (SEQ ID NO:7), 5'-GCTGCTGATGGTGA-GAGTGA (SEQ ID NO:8)

(2) Fluorescence in situ hybridization (FISH)

35 FISH analysis was conducted with probes specific to human chromosomes #2, 4, 14 and 22 (CHROMOSOME PAINTING SYSTEM, Cambio Ltd.) in accordance with the method described in Matsubara et al., "FISH Experiment Protocol", published by Shujunsha, 1994.

For example, at least one clone retaining chromosome #2 was obtained in 10 groups out of 26 groups (745 clones). 40 Among them, only 5 clones were positive to all the used primers specific to chromosome #2. FISH analysis was conducted with these clones. FISH analysis was conducted with probes specific to human chromosomes #2 (CHROMOSOME PAINTING SYSTEM, Cambio Ltd.) in accordance with the method described in Matsubara et al., "FISH Experiment Protocol", published by Shujunsha, 1994. In the cells positive to all the primers, an intact form of human chromosome #2 was observed. In some of the clones positive to part of the primers, an independent chromosome 45 smaller than human chromosome #2 was observed or a cell having a chromosome in a form fusing with chromosomes other than human chromosome #2 was observed (Fig. 1). In Fig. 1, the names of the clones are shown in the horizontal line and the primers used in the PCR are shown in the left longitudinal line. ● shows positive clones and X shows negative clones. The forms of human chromosome #2 observed by FISH are shown in the bottom line. No description means no performance of experiment.

50 A9 cells retaining human chromosomes #4, 14 and 22 were obtained by the same procedure.

Example 2

Transfer of human chromosome #22 into mouse ES cells by microcell fusion

55 The mouse A9 cell clones retaining human chromosome #22 (hereinafter referred to as "A9/#22") from Example 1 were used as chromosome donor cells. Mouse ES cell line E14 (obtained from Martin L. Hooper; Hooper et al., Nature, 326:292, 1987) was used as a chromosome recipient cell. E14 cells were cultured in accordance with the method described in Aizawa Shinichi, "Biomanual Series 8, Gene Targeting", published by Yodosha, 1995 and G418 resistant

STO cell line (obtained from Prof. Kondo Hisato, Osaka University) treated with mitomycin C (Sigma) was used as a feeder cell. In the first step, microcells were prepared from about 10^8 cells of A9/#22 in accordance with the method reported by Shimizu et al. "Cell Technology Handbook", published by Yodosha, 1992. The total amount of the resulting microcells were suspended in 5 ml of DMEM. About 10^7 cells of E14 were dispersed with trypsin and washed three times with DMEM and suspended in 5 ml of DMEM. The cells were then mixed with the microcells and the mixture was centrifuged at 1,250 rpm for 10 minutes to remove the supernatant. The precipitate was dispersed by tapping and 0.5 ml of a PEG solution (1:1.4) [5 g of PEG 1000 (Wako Pure Chemicals Co., Ltd.) and 1 ml of DMSO (Sigma) as dissolved in 6 ml of DMEM] was added. The mixture was left to stand at room temperature for 1 minute and 30 seconds and 10 ml of DMEM was added slowly. Immediately thereafter, the resulting mixture was centrifuged at 1,250 rpm for 10 minutes to remove the supernatant. The precipitate was suspended in 30 ml of a medium for ES cells and inoculated into 3 tissue culture plastic plates (Corning) of 100 mm in diameter into which feeder cells were inoculated. After 24 hours, the medium was replaced with a medium supplemented with 300 µg/ml of G418 (GENETICIN, Sigma) and medium replacements were thereafter conducted daily. Drug resistant colonies appeared in 1 week to 10 days. The frequency of appearance was 0.5 per 10^7 of E14 cells. The colonies were picked up and grown. The cells were suspended in a storage medium (a medium for ES cells + 10% DMSO (Sigma)) at a concentration of 5×10^6 cells per ml and stored frozen at -80°C. At the same time, genomic DNA was prepared from 10^6 - 10^7 cells of each drug resistant clone with a Puregene DNA Isolation Kit (Gentra System Co.).

Human chromosome #22 was fragmented by irradiating the microcells with γ rays (Koi et al., Science, 260:361, 1993). The microcells obtained from about 10^8 cells of A9/#22 were suspended in 5 ml of DMEM and irradiated with γ rays of 60 Gy on ice with a Gammacell 40 (Canadian Atomic Energy Public Corporation) at 1.2 Gy/min for 50 minutes. The fusion of γ ray-irradiated microcells and the selection of drug resistant clones were conducted by the same procedure as in the case of the unirradiated microcells. As a result, the frequency of the appearance of the drug resistant clones was 1.7 per 10^7 of E14 cells. The drug resistant clones were stored frozen and DNA was prepared from the clones by the same procedure as in the case of the unirradiated microcells.

The retention of the transferred chromosomes in the unirradiated microcell-transferred drug resistant clones E14/#22-9 and E14/#22-10, and in the γ ray-irradiated microcell-transferred drug resistant clones E14/#22-14 and E14/#22-25 was confirmed by the methods described in (1)-(3) below.

(1) PCR analysis (Fig. 2)

The presence of the gene on human chromosome #22 (Genetic Maps, supra) and polymorphic markers (Polymorphic STS Primer Pairs: D22S315, D22S275, D22S278, D22S272 and D22S274, BIOS Laboratories, Inc.; Nature 359:794, 1992) was detected by a PCR method using the genomic DNA of the drug resistant clone as a template. The sequences of oligonucleotide primers for the genes prepared on the basis of nucleotide sequences obtained from data bases such as GenBank, EMBL and the like are described below.

PVALB (parvalbumin) : 5'-TGGTGGCTGAAAGCTAAGAA (SEQ ID NO:9), 5'-CCAGAAGAATGGTGTCATTA (SEQ ID NO:10)

MB (myoglobin) : 5'-TCCAGGTTCTGCAGAGCAAG (SEQ ID NO:11), 5'-TGTAGTTGGAGGCCATGTCC (SEQ ID NO:12)

DIA1 (cytochrome b-5 reductase) : 5'-CCCCACCCATGATCCAGTAC (SEQ ID NO:13), 5'-GCCCTCAGAACGACGAAG-CAG (SEQ ID NO:14)

Ig λ (immunoglobulin lambda) : 5'-GAGAGTTGCAGAAGGGGTGACT (SEQ ID NO:15), 5'-GGAGACCACCAAAC-CCTCCAAA (SEQ ID NO:16)

ARSA (arylsulfatase A) : 5'-GGCTATGGGGACCTGGGCTG (SEQ ID NO:17), 5'-CAGAGACACAGGCACGTAGAAG (SEQ ID NO:18)

PCR amplification (Innis et al., supra) was conducted by using about 0.1 µg of the genomic DNA as a template with the above 10 kinds of the primers. As a result, amplification products having expected lengths were detected with all the primers in the case of the two unirradiated clones and with part of the primers in the case of the γ ray-irradiated two clones. The results are shown in Fig. 2. In Fig. 2, a schematic chromosome map based on the G bands of human chromosome #22 and the location of some markers on bands are shown at the left side (O'Brien, GENETIC MAPS, 6th edition, BOOK 5, etc.). The arrangement of the genetic and polymorphic markers shows approximate positional relationships on the basis of the presently available information (Science, HUMAN GENETIC MAP, 1994; Nature Genetics, 7:22, 1994; Nature 359:794, 1992, etc.) and the order is not necessarily correct. With respect to four kinds of the G418 resistant E14 cell clones, the markers for which the expected amplification products were detected by PCR are shown by ■ and the markers for which the expected amplification products were not detected are shown by □. The results of the observation by FISH analysis are shown at the bottom side. A9/#22 is a chromosome donor cell.

(2) Southern blot analysis

South rn blot analysis of about 2 µg of the genomic DNA digested with restriction enzyme BgIII (TAKARA SHUZO CO., LTD.) was conducted by using human specific repeated sequence L1 (10^4 - 10^5 copies were present per haploid genome, obtained from RIKEN DNA Bank; Nucleic acids research, 13;7813, 1985; pUK19A derived EcoRI-BamHI fragment of 1.4 kb) as a probe in accordance with the method described in Ausubel et al., Current Protocols in Molecular Biology, John Wiley & Sons, Inc., 1994. As a result, a large number of bands hybridized with the human L1 sequence were detected in DNA of each drug resistant clone. With respect to the unirradiated 2 clones, their patterns and the quantitative ratio of human chromosomal DNA to mouse genomic DNA which could be presumed from the density of the respective bands were the same as those of A9/#22. The total signal intensity of the bands of the γ -ray irradiated clones correlated with the degree of the deletion confirmed by the PCR analysis, as compared with that of A9/#22.

(3) Fluorescence in situ hybridization (FISH)

FISH analysis was conducted with probes specific to human chromosomes #22 (CHROMOSOME PAINTING SYSTEM, Cambio Ltd.) in accordance with the method described in Matsubara et al., "FISH Experiment Protocol", published by Shujunsha, 1994. As a result, in almost all of the observed metaphase spreads, human chromosome #22 was detected in the form of translocation to the mouse chromosome with respect to E14/#22-9 and in the form of an independent chromosome with respect to the three other clones.

The results of the above experiments demonstrate that the obtained G418 resistant clones E14/#22-9 and E14/#22-10 retained all or most part of human chromosome #22 whereas the clones E14/#22-14 and E14/#22-25 retained partial fragments of human chromosome #22.

Example 3

Production of chimeric mice from the ES cells retaining human chromosome #22

General procedures for obtaining mouse embryos, cultivation, injection of the ES cells into the embryos, transplantation to the uteri of foster mothers were carried out in accordance with the method described in Aizawa Shinichi, "Biomanual Series 8, Gene Targeting", published by Yodoshia, 1995. The cells in a frozen stock of the G418 resistant ES clone E14/#22-9 which was confirmed to retain human chromosome #22 were thawed, started to culture and injected into blastocyst-stage embryos obtained by mating a C57BL/6XC3H F1 female mouse (CREA JAPAN, INC.) with a C3H male mouse (CREA JAPAN, INC.); the injection rate was 10-15 cells per embryo. Two and half days after a foster mother [ICR or MCH(ICR)] mouse (CREA JAPAN, INC) was subjected to a pseudopregnant treatment, about ten of the ES cell-injected embryos were transplanted to each side of the uterus of the foster mother. The results are shown in Table 1.

Table 1

Production of chimeric mice from the ES cells retaining human chromosome #22 (fragments)							
ES cell clone/human chromosome	G418 resistant clone No.	Number of ES cell-injected blastocyst stage embryos	Number of off-spring mice	Number of chimeric mice	Contribution to coat color		
					<10%	10-30%	30%<
E14/#22	9	166	29	16	7	3	6

As a result of the transplantation of a total 166 of injected embryos, 29 offspring mice were born. Chimerism in the offsprings can be determined by the extent of E14 cell-derived pale gray coat color in the host embryo-derived agouti coat color (dark brown). Out of the 29 offsprings, 16 mice were recognized to have partial pale gray coat color, indicating the contribution of the E14 cells. The maximum contribution was about 40% in K22-22.

These results show that the mouse ES cell clone E14/#22-9 retaining human chromosome #22 maintains the ability to produce chimera, that is, the ability to differentiate into normal tissues of mouse.

Example 4

Confirmation of retention of human chromosomal DNA in various tissues of the chimeric mice derived from the ES cells retaining human chromosome #22

In addition to the determination of coat color in Example 3, the retention of the transferred chromosome was confirmed by PCR analysis using a template genomic DNA prepared from the tail of the chimeric mouse. The tail was obtained from the chimeric mouse at least 3 weeks old in accordance with the method described in Motoya Katsuki, "Development Technology Experiment Manual", published by Kodansha Scientific, 1987. Genomic DNA was extracted from the tail with a Puregene DNA Isolation Kit. Out of the polymorphic primers used in Example 2, PVALB and D22S278 were used, with the extracted genomic DNA as a template, to confirm the amplification products. The analysis was conducted with 10 of the mice in which the contribution to coat color was observed. As a result, the products of amplification with at least either of the primers were detected in all the mice.

Southern blot analysis was conducted in the same manner as in Example 2 by using human L1 sequence as a probe with 2 µg of the genomic DNA derived from the tails of the 6 chimeric mice and one non-chimeric mouse. As a result, the presence of a large number of human L1 sequence was observed in all the chimeric mice and their patterns were similar to those of E14/#22-9. The quantitative ratio to mouse genome was about 10% at maximum (Fig. 3). In Fig. 3, 2 µg of genomic DNA digested with BglII was used in each lane. Human L1 sequence labeled with ³²P was used as a probe and signals were detected with Image Analyzer BAS2000 (Fuji Photo Film Co., Ltd.). The lanes represent the genomic DNA derived from the tails of the chimeric mice (K22-6, 7, 8, 9, 10, 11 and 12; 9 is the non-chimeric mouse) and control DNA (C which is a mixture of E14/#22-9 genomic DNA and E14 genomic DNA at a weight ratio of 1:9) as counted from the right. The DNA molecular weights are shown at the left side and chimerism in the chimeric mice at the right side (-: 0%, +: <10%, and ++: 10-30%).

With respect to the chimeric mouse (K22-7) having about 5% contribution to coat color, genomic DNA was obtained from the brain, liver, muscle, heart, spleen, thymus, ovary and kidney with an ISOGEN (Nippon Gene Co.). For each tissue, PCR analysis was conducted with MB and D1A1 selected from the primers for the genes used in Example 2. As a result, both primers gave expected amplification products in all the tissues. The results of PCR analysis using D1A1 primer are shown in Fig. 4. The PCR products were electrophoresed on a 2% agarose gel and stained with ethidium bromide for detection. The lanes in Fig. 4 represent the following from the left: B, brain; L, liver; SM, skeletal muscle; H, heart; Sp, Spleen; Th, thymus; Ov, ovary; K, kidney; nc, non-chimeric mouse tail-derived DNA (negative control); pc, human fibroblast cell (HFL-1) DNA (positive control).

These results show that E14/#22-9 contributed to various normal tissues in the mouse and that it retained human chromosome #22.

Example 5

Expression of the human genes in the chimeric mouse derived from the ES cell retaining human chromosome #22

The tail of the mouse (K22-7) having about 5% contribution to coat color was frozen with liquid nitrogen and then disrupted for use as a sample for confirming the expression of the human genes. The sample was a mixture of tissues such as skin, bones, muscles, blood and the like. Total RNA was extracted from the sample with an ISOGEN (Nippon Gene Co.) and used in an RT-PCR method to detect mRNAs of human myoglobin (MB) and human cytochrome b5 reductase (D1A1). The RT-PCR was performed in accordance with the method described in Innis et al., "PCR Experiment Manual", published by HBJ Publication Office, 1991. Random hexamer oligonucleotides (final concentration: 100 pmol, TAKARA SHUZO CO., LTD.) were used as primers for reverse transcription and Super Script (BRL Co.) as reverse transcriptase. The following primers were used for amplification using cDNA as a template.
 MB: 5' -TTAAGGGTCACCCAGAGACT (SEQ ID NO:19), 5'-TGTAGTTGGAGGCCATGTCC (SEQ ID NO:20)
 D1A1: 5'-CAAAAGTCCAACCTATCA (SEQ ID NO:21), 5'-GCCCTCAGAAGACGAAGCAG (SEQ ID NO:22)

As a result, amplification products specific to mRNAs of both genes were detected (Fig. 5). The RT-PCR products were electrophoresed on a 2% agarose gel and stained with ethidium bromide for detection. In Fig. 5, M is a marker (HindIII digested λ DNA + HaeIII digested φ X174DNA, TAKARA SHUZO CO., LTD.); MB, human myoglobin; D1A1, human cytochrome b5 reductase; and WT, a wild-type C3H mouse.

With respect to the same individual (K22-7), total RNA was extracted from the brain, heart, thymus, liver, spleen, kidney, ovary and skeletal muscle with an ISOGEN and RT-PCR was performed on each organ with the above two primers. As a result, expected products of amplification with D1A1 were observed in all the organs and those with MB were observed only in the heart and skeletal muscle (Fig. 6). Myoglobin is known to be expressed specifically in muscle cells (Bassel-Duby et al., MCB, 12:5024, 1992). Hence, the above results show that the gene on the transferred human chromosome can be subjected to the normal tissue-specific regulation in the mouse. The PCR products were electrophoresed on a 2% agarose gel and stained with ethidium bromide for detection. In Fig. 6, the lanes represent the

following from the left: B, brain; H, heart; Th, thymus; L, liver; Sp, spleen; K, kidney; Ov, ovary; SM, skeletal muscle; and M, marker (supra). The lower band observed in the results of MB are believed to represent non-specific products.

These results show that the transferred human chromosome #22 can function in normal tissues of the chimeric mice.

5

Example 6

Transfer of human chromosome #4 or fragments thereof into ES cells

10 The mouse A9 cell clone retaining human chromosome #4 (hereinafter referred to as "A9/#4") from Example 1 was used as a chromosome donor cell. Mouse ES cell line E14 (see Example 2) was used as a chromosome recipient cell. The microcell fusion and the selection of G418 resistant clones were conducted by the same procedures as in Example 2. The frequency of the appearance of the drug resistant clones was 1-2 per 10^7 of E14 cells. The drug resistant clones were stored frozen and genomic DNA were prepared by the same procedures as in Example 2. The retention of the 15 transferred human chromosome #4 or fragments thereof in the drug resistant clones E14/#4-4, E14/#4-7 and E14/#4-11 was confirmed by the methods described in (1)-(3) below.

(1) PCR analysis (Fig. 7)

20 The presence of the gene on human chromosome #4 (O'Brien, Genetic Maps, 6th edition, Book 5, Cold Spring Harbor Laboratory Press, 1993) and polymorphic markers (Polymorphic STS Primer Pairs: D4S395, D4S412, D4S422, D4S413, D4S418, D4S426 and F11, BIOS Laboratories, Inc.; Nature 359:794, 1992) was detected by a PCR method. The sequences of oligonucleotide primers for the genes prepared on the basis of nucleotide sequences obtained from data bases such as GenBank, EMBL and the like will be described below.

25 HD (huntington disease) : 5'-TCGTTCCCTGTCGAGGATGAA (SEQ ID NO:23), 5'-TCACTCCGAAGCTGCCTTTC (SEQ ID NO:24)

IL-2 (interleukin-2) : 5'-ATGTACAGGATGCAACTCCTG (SEQ ID NO:25), 5'-TCATCTGTAATCCAGCAGT (SEQ ID NO:26)

KIT (c-kit) : 5'-GATCCCCTCGCAGCTACCGC (SEQ ID NO:27), 5'-TTCGCCGAGTAGTCGCACGG (SEQ ID NO:28)

30 FABP2 (fatty acid binding protein 2, intestinal), 5'-GATGAACTAGTCCAGGT GAGTT (SEQ ID NO:29), 5'-CCTTTT-GGCTTCTACTCCTTCA (SEQ ID NO:30)

PCR amplification was conducted with the above 11 kinds of the primers. As a result, the amplification products having expected lengths were detected with all or part of the primers in all the three clones. In the E14/#4-4 and E14/#4-7 clones, the deletion of partial regions was observed. The results are shown in Fig. 7. In Fig. 7, a schematic chromosome map based on the G bands of human chromosome #4 and the location of some markers on bands are shown at the left side (see Example 2). The arrangement of the genetic and polymorphic markers shows approximate positional relationships on the basis of the presently available information (see Example 2) and the order is not necessarily correct. With respect to the three kinds of the G418 resistant E14 cell clones, the markers for which the expected amplification products were detected are shown by ■ and the markers for which the expected amplification products were not 40 detected are shown by □. The results of the observation by FISH analysis are shown at the lower side. A9/#4 is a chromosome donor cell.

(2) Southern blot analysis (Fig. 8)

45 Southern blot analysis was conducted by the same procedure as in Example 2 using human L1 sequence as a probe with genomic DNAs obtained from E14/#4-4 and E14/#4-7. As a result, a large number of bands hybridized with the human L1 sequence were detected in DNAs of both drug resistant clones. The total signal intensity correlated with the degree of the deletion confirmed by the PCR analysis, as compared with that of A9/#4. In Fig. 8, 2 µg of genomic DNA digested with BglII was used in each lane. Human L1 sequence labeled with 32 P was used as a probe and the 50 signals were detected with an Image Analyzer (BAS 2000, Fuji Photo Film Co., Ltd.). In Fig. 8, the lanes represent the following as counted from the left: 1, A9/#4 (chromosome donor cell); 2, A9/#4+A9 (1:2); 3, A9/#4+A9 (1:9); 4, A9; 5, E14/#4-7; and 6, E14/#4-4. Lanes 2 and 3 represent mixtures of two kinds of DNAs at the ratios shown in parentheses. The molecular weights of DNAs are shown at the left side.

55 (3) Fluorescence in situ hybridization (FISH)

FISH analysis was conducted with probes specific to human chromosomes #4 (CHROMOSOME PAINTING SYSTEM, Cambio Ltd.) by the same procedure as in Example 2. As a result, in almost all of the observed metaphase spreads of the three clones used, human chromosome #4 or partial fragments thereof were detected in the form of

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translocation to the mouse chromosome with respect to E14/#4-4 and in the form of an independent chromosome with respect to the two other clones. The relative sizes of the observed human chromosome were consistent with those presumed from the results of the PCR analysis.

5 The results of the above experiments demonstrate that the obtained G418 resistant clones retained the whole human chromosome #4 or partial fragments thereof.

Example 7

10 Production of chimeric mice from the ES cells retaining human chromosome #4 fragments

15 The cells in frozen stocks of the G418 resistant ES cell clones E14/#4-4 and E14/#4-7 which were confirmed to retain partial fragments of human chromosome #4 were thawed, started to culture, and injected into blastocyst stage embryos obtained by the same method as in Example 3; the injection rate was 10-15 cells per embryo. Two and half days after a foster mother [ICR or MCH(ICR)] mouse (CREA JAPAN, INC.) was subjected to a pseudopregnant treatment, about ten of the ES cell-injected embryos were transplanted to each side of the uterus of the foster mother. The results are shown in Table 2.

Table 2

Production of chimeric mice from the E14 cell clones retaining human chromosome #4 (fragments)							
ES cell clone/human chromosome	G418 resistant clone No.	Number of ES cell-injected blastocyst stage embryos	Number of offspring mice	Number of chimeric mice	Contribution to coat color		
					<10%	10-30%	30%<
E14/#4	4	160	8	5	5	-	-
	7	80	5	2	1	1	-

30 As a result of the transplantation of a total of 240 injected embryos, 13 offspring mice were born. Chimerism in the offsprings can be determined by the extent of E14 cell-derived pale gray coat color in the host embryo-derived agouti coat color (dark brown). Out of the 13 offsprings, 7 mice were recognized to have partial pale gray coat color, indicating the contribution of the E14 cells. The maximum contribution was about 15% in one individual derived from E14/#4-7.

35 These results show that the mouse ES cell clones E14/#4-4 and E14/#4-7 which retain fragments of human chromosome #4 maintain the ability to produce chimera, that is, the ability to differentiate into normal tissues of mouse.

40 Example 8

Confirmation of retention of human chromosomal DNA in the chimeric mice derived from the ES cells retaining partial fragments of human chromosome #4 and expression of the G418 resistance gene

45 (1) PCR analysis

Using the chimeric mice produced in Example 7, genomic DNAs were prepared from the tails of one individual derived from E14/#4-7 (K#4-7-1: about 5% chimerism) and one individual derived from E14/#4-4 (K#4-4-41: about 5% chimerism) by the same procedure as in Example 4. These DNAs were used as templates to conduct PCR analysis 50 using polymorphic marker F11 for chromosome #4 analysis (see Example 6) which was detected in E14/#4-7 and E14/#4-4. As a result, expected amplification products were detected in both mice.

(2) Southern analysis (Fig. 9)

55 Southern analysis was conducted in the same manner as in Example 2 by using human L1 sequence as a probe with 2 µg of the genomic DNA derived from the tail of one individual derived from E14/#4-7 (K#4-7-1: about 5% chimerism). As a result, the presence of a large number of human L1 sequence was observed and their patterns were similar to those of E14/#4-7. The quantitative ratio to mouse genome was about 10% of that of E14/#4-7 at maximum. In Fig. 9, 2 µg of genomic DNA digested with BglII was used in each lane. Human L1 sequence labeled with ³²P was used as

a probe and signals were detected with Image Analyzer BAS2000 (Fuji Photo Film Co., Ltd.). The molecular weights of DNAs are shown at the left side. The lanes represent the following as counted from the left: 1, K#4-7-1; 2, blank; and 3, E14/#4-7.

5 (3) Test on the tail-derived fibroblast cells for G418 resistance

Fibroblast cells were prepared from the tails of one individual derived from E14/#4-7 (K#4-7-1: about 5% chimerism) and one individual derived from E14/#4-4 (K#4-4-41: about 5% chimerism). In the same procedure as in Example 4, the tail of each mouse was cut at a length of 5-10 mm and washed several times with PBS/1mM EDTA, followed by 10 notching of the tail with a knife. The outer skin layer was removed and the inner tissues were cut into fine pieces. The fine pieces of tissues were transferred into a tube containing 5 ml of PBS/1 mM EDTA and left to stand for 30 minutes to 1 hour at room temperature. Subsequently, the supernatant was removed leaving a 1 ml portion of the PBS/EDTA behind, and 1 ml of 0.25% trypsin/PBS was added. The tissues were dispersed thoroughly by tapping or pipetting at room temperature for 5-10 minutes. After centrifugation at 1,000 rpm for 10 minutes, the precipitate was suspended in 15 2 ml of DMEM (10% FCS) and inoculated into a 35 mm plate. After cultivation for 7-10 days, the cells were treated with trypsin and about 10^4 cells per plate were inoculated into two 35 mm plates. G418 was added to the medium in one plate at a final concentration of 400 μ g/ml. The cells were cultured for 5-7 days and the appearance of viable cells in each plate were examined. Under these conditions, 100% of the wild-type ICR mouse-derived fibroblast cells were killed in the presence of G418. As a result, G418 resistant fibroblast cells was present in both mice.

20 These results show that E14/#4-7 and E14/#4-4 contributed to various normal tissues in the mouse and that they retained partial fragments of human chromosome #4.

Example 9

25 Transfer of human chromosome #14 or fragments thereof into mouse ES cells

The mouse A9 cell clone retaining human chromosome #14 (hereinafter referred to as "A9/#14") from Example 1 was used as a chromosome donor cell. Mouse ES cell line TT2 (purchased from Lifetech Oriental Co., Yagi et al., Analytical Biochem., 214:70, 1993) was used as a chromosome recipient cell. The TT2 cells were cultured in accordance 30 with the method described in Aizawa Shinichi, "Biomanual Series 8, Gene Targeting", published by Yodosha, 1995 and G418 resistant primary culture cells (purchased from Lifetech Oriental Co.) treated with mitomycin C (Sigma) were used as feeder cells. The microcell fusion and the selection of G418 resistant clones were conducted by the same procedures as in Example 2. The frequency of the appearance of the drug resistant clones was 3-6 per 10^7 of TT2 cells. The drug resistant clones were stored frozen and genomic DNA was prepared by the same procedures as in Example 2.

35 Human chromosome #14 was fragmented by irradiating the microcells with γ -rays (Koi et al., Science, 260:361, 1993). The microcells obtained from about 10^8 cells of A9/#14 were suspended in 5 ml of DMEM and irradiated with γ -rays of 30 Gy on ice with a Gammacell 40 (Canadian Atomic Energy Public Corporation) at 1.2 Gy/min for 25 minutes. The fusion of γ ray-irradiated microcells and the selection of drug resistant clones were conducted by the same procedure 40 as in the case of the unirradiated microcells. As a result, the frequency of the appearance of the drug resistant clones was 3 per 10^7 of TT2 cells. The drug resistant clones were frozen stored and DNA was prepared by the same procedure as in Example 2.

The retention of human chromosome #14 or partial fragments thereof in the unirradiated microcell-transferred G418 resistant clones 1-4 and 1-5, and in the G418 resistant clones 3-1 and 3-2 (a total of 4 clones) into which the γ -ray-irradiated microcell was transferred was confirmed by the methods described in (1) and (2) below.

45

(1) PCR analysis (Fig. 10)

The presence of the gene on human chromosome #14 (O'Brien, Genetic Maps, 6th edition, Book 5, Cold Spring 50 Harbor Laboratory Press, 1993) and polymorphic markers (Polymorphic STS Primer Pairs: D14S43, D14S51, D14S62, D14S65, D14S66, D14S67, D14S72, D14S75, D14S78, D14S81 and PCI, BIOS Laboratories, Inc.; Nature 359:794, 1992; Nature Genetics, 7:22, 1994) was detected by a PCR method using genomic DNA of the drug resistant clone as a template. The sequences of oligonucleotide primers for the genes prepared on the basis of nucleotide sequences obtained from data bases such as GenBank, EMBL and the like are described below.

NP (nucleoside phosphorylase) : 5'-ATAGAGGGTACCCACTCTGG (SEQ ID NO:31), 5'-AACCAAGGTAGGTTGATATGG (SEQ ID NO:32)

TCRA (T-cell receptor alpha) : 5'-AAGTTCCCTGTGATGTCAAGC (SEQ ID NO:33), 5'-TCATGAGCAGATTAAACCCG (SEQ ID NO:34)

MYH6 (myosin heavy chain cardiac) : 5'-TGTGAAGGAGGACCAGGTGT (SEQ ID NO:35), 5'-TGTAGGGGTTGACAGTGACA (SEQ ID NO:36)

IGA2 (immunoglobulin alpha-2 constant) : 5'-CTGAGAGATGCCTCTGGTGC (SEQ ID NO:37), 5'-GGCGGTTAGT-GGGGTCTTCA (SEQ ID NO:38)

IGG1 (immunoglobulin gamma-1 constant) : 5'-GGTGTCTGGAACTCAGGCG (SEQ ID NO:39), 5'-CTGGTGCAG-GACGGTGAGGA (SEQ ID NO:40)

5 IGM (immunoglobulin mu constant) : 5'-GCATCCTGACCGTGTCCGAA (SEQ ID NO:41), 5'-GGGTCAGTAGCAGGT-GCCAG (SEQ ID NO:42)

IGVH3 (immunoglobulin heavy variable-3) : 5'-AGTGAGATAAGCAGTGGATG (SEQ ID NO:43), 5'-GTTGTGCTACTC-CCATCACT (SEQ ID NO:44)

10 PCR amplification was conducted using the genomic DNAs of the 4 drug resistant clones as templates with the above 18 kinds of the primers by the same procedure as in Example 2. As a result, expected amplification products were detected with all or part of the primers. In the drug resistant clones 3-1 and 3-2 obtained by using the γ -ray irradiated microcells, a tendency for the deletion of partial regions of chromosome #14 was observed. In the case where the unirradiated microcells were used, deletion was observed as in the case of the 1-4 clone. The results are shown in Fig. 10. In Fig. 10, a schematic chromosome map based on the G bands of human chromosome #14 and the location of some markers on bands are shown at the left side (see Example 2). The arrangement of the genetic and polymorphic markers shows approximate positional relationships on the basis of the presently available information (see Example 2) and the order is not necessarily correct. With respect to four kinds of the G418 resistant TT2 cell clones, the markers for which the expected amplification products were detected are shown by ■ and the markers for which the expected amplification products were not detected are shown by □. A9/#14 is a chromosome donor cell. The results of Example 11 (1) are shown at the right side.

(2) Fluorescence in situ hybridization (FISH)

25 FISH analysis was conducted with probes specific to human chromosomes #14 (CHROMOSOME PAINTING SYSTEM, Cambio Ltd.) in accordance with the method described in Matsubara et al., "FISH Experiment Protocol", published by Shujunsha, 1994. As a result, in almost all of the observed metaphase spreads of all the 4 clones, human chromosome #14 or partial fragments thereof were detected in the form of an independent chromosome. The relative sizes of the observed human chromosome were consistent with those presumed from the results of the PCR analysis.

30 The results of the above experiments demonstrate that the obtained G418 resistant clones 1-4, 1-5, 3-1 and 3-2 retained the whole or partial fragments of human chromosome #14.

Example 10

Production of chimeric mice from the ES cells retaining human chromosome #14 or fragments thereof

35 The cells in the frozen stocks of four G418 resistant ES cell clones (1-4, 3-1, 3-2 and 1-5) that were prepared in Example 9 and which were confirmed to retain human chromosome #14 or fragments thereof were thawed, started to culture and injected into 8-cell stage embryos obtained by mating [ICR or MCH(ICR)] male and female mice (CREA JAPAN, INC.); the injection rate was 8-10 cells per embryo. The embryos were cultured in an ES medium overnight to develop to blastocysts. Two and half days after a foster mother ICR mouse (CREA JAPAN, INC.) was subjected to a pseudopregnant treatment, about ten of the injected embryos were transplanted to each side of the uterus of the foster mother. The results are shown in Table 3.

Table 3

Production of chimeric mice from the TT2 cell clones retaining human chromosome #14 (fragments)							
ES cell clone/human chromosome	G418 resistant clone No.	Number of ES cell-injected 8-cell stage embryos	Number of off-spring mice	Number of chimeric mice	Contribution to coat color		
TT2/#14	1-4	98	20	1	-	-	1
	1-5	110	14	2	1	-	1
	3-1	103	11	2	1	1	-
	3-2	183	19	3	-	2	1

As a result of the transplantation of a total of 494 injected embryos, 64 offspring mice were born. Chimerism in the offsprings can be determined by the extent of TT2 cell-derived agouti coat color (dark brown) in the host embryo-derived albinos coat color. Out of the 64 produced offsprings, 8 mice were recognized to have partial agouti coat color, indicating the contribution of the ES cells. The maximum contribution was about 80% in one individual derived from 1-4.

These results show that the G418 resistant ES cell clones (1-4, 1-5, 3-1 and 3-2) retaining human chromosome #14 or fragments thereof maintain the ability to produce chimera, that is, the ability to differentiate into normal tissues of mouse.

10 Example 11

Confirmation of retention of human chromosome #14 fragment DNA in the chimeric mice derived from the ES cells retaining human chromosome #14 fragments

15 The retention of human chromosome #14 partial fragments in the chimeric mice obtained in Example 10 was confirmed by the methods described in (1)-(3) below.

(1) PCR analysis using DNAs derived from various tissues

20 Genomic DNA was extracted from the tail of one individual derived from 3-1 (K3-1-1: about 25% chimerism) by the same procedure as in Example 4. The DNA was used as a template to conduct PCR analysis using all of the 14 primers for chromosome #14 analysis which were detected in 3-1. As a result, expected amplification products were detected with all the 14 primers. (Fig. 10)

With respect to the same individual (K3-1-1), genomic DNA was obtained from the brain, kidney, spleen, heart, liver 25 and thymus with a Puregene DNA Isolation Kit. For each tissue, PCR analysis was conducted with IGM primers (see Example 9). As a result, expected amplification products were detected in all the tissues (Fig. 11). The PCR products were electrophoresed on a 2% agarose gel and stained with ethidium bromide for detection. In Fig. 11, the lanes represent the following as counted from the left: B, brain; K, kidney; Sp, Spleen; H, heart; L, liver; Th, thymus; pc, human fibroblast cell (HFL-1) DNA (positive control); nc, non-chimeric mouse tail DNA (negative control); and M, marker (Hin-dIII digested λ DNA + HaeIII digested φ X174 DNA, TAKARA SHUZO CO., LTD.).

(2) Test on the tail-derived fibroblast cells for G418 resistance

Fibroblast cells were prepared from the tails of two individuals derived from 3-2 (K3-2-1: about 25% chimerism, and 35 K3-2-3: about 50% chimerism) and one individual derived from 1-4 (K1-4-1: about 80% chimerism). In the same procedure as in Example 4, the tail of each chimeric mouse of 3-6 weeks was cut at a length of 5-10 mm and washed several times with PBS/1mM EDTA, followed by notching of the tail with a knife. The outer layer was removed and the inner tissues were cut into fine pieces. The fine pieces of tissues were transferred into a tube containing 5 ml of PBS/1 mM EDTA and left to stand for 30 minutes to 1 hour at room temperature. Subsequently, the supernatant was removed leaving a 1 ml portion of the PBS/EDTA behind, and 1 ml of 0.25% trypsin/PBS was added. The tissues were dispersed thoroughly by tapping or pipetting at room temperature for 5-10 minutes. After centrifugation at 1,000 rpm for 10 minutes, the precipitate was suspended in 2 ml of DMEM (10% FCS) and inoculated into a 35 mm plate. After cultivation for 7-10 days, the cells were treated with trypsin and about 10^4 cells per plate were inoculated into four 35 mm plates. 40 G418 was added to the medium in two of the plates at a final concentration of 400 µg/ml. The cells were cultured for 5-7 days and the viable cells in each plate were counted. Under these conditions, 100% of the wild-type ICR mouse-derived fibroblast cells were killed in the presence of G418. Assuming the same growth rate of the G418 resistant fibroblast in the non-selective and selective media, the ratio of the viable cells in the selective medium to those in the non-selective medium is believed to reflect the contribution in the fibroblast cell populations of the G418 resistant ES cell-derived fibroblast. As a result, the presence of G418 resistant fibroblast cells was observed in all the three individuals 45 as shown in Fig. 12. In Fig. 12, % resistance is an average of 2 pairs of the selective/non-selective 35 mm plates for each mouse. ICR refers to the wild-type ICR mice.

(3) FISH analysis of the tail-derived G418 resistant fibroblast cells

55 FISH analysis of the K3-2-3 and K1-4-1 derived G418 resistant fibroblast cells obtained in (2) was conducted by the same procedure as in Example 2. Total human DNA extracted from the HFL-1 cells (Example 1) was labeled with FITC so that it could be used as a probe (Matsubara et al., "FISH Experimental Protocol", published by Shujunsha, 1994). As a result, in almost all of the observed metaphase spreads of the both individuals, partial fragments of the human chromosome in independent forms were observed.

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These results show that the TT2 cell clones retaining fragments of human chromosome #14 contributed to various normal tissues in the mouse individuals and that they retained partial fragments of human chromosome #14.

Example 12

5

Transfer of partial fragments of human chromosome #2 into ES cells

10

The mouse A9 cell W23 retaining a human chromosome #2 fragment (hereinafter referred to as "A9/#2 W23") from Example 1 was used as a chromosome donor cell. Mouse ES cell line TT2 (see Example 9) was used as a chromosome recipient cell. The microcell fusion and the selection of G418 resistant clones were conducted by the same procedures as in Example 2. The frequency of the appearance of the drug resistant clones was 1-3 per 10^7 of TT2 cells. The drug resistant clones were stored frozen and genomic DNA was prepared by the same procedures as in Example 2. The retention of partial fragments of human chromosome #2 in drug resistant clones 5-1, 5-2 and 5-3 was confirmed by the methods described in (1) and (2) below.

15

(1) PCR analysis

20

The presence of Ck and FABP1 that are the genes on human chromosome #2 (Genetic Maps, supra) and which were detected in the chromosome donor cell A9/#2 W23 was detected by a PCR method.

25

As a result of PCR amplification using each primer, expected amplification products were detected with both primers in all of the 3 clones.

(2) Fluorescence in situ hybridization (FISH)

25

FISH analysis was conducted with probes specific to human chromosome #2 (CHROMOSOME PAINTING SYSTEM, Cambio Ltd.) by the same method as in Example 2. As a result, in almost all of the observed metaphase spreads of the 3 clones, partial fragments of human chromosome #2 in the form of independent chromosomes were detected. The sizes of the observed human chromosome were the same as those observed in A9/#2 W23.

30

The results of the above experiments demonstrate that the obtained G418 resistant clones retained partial fragments of human chromosome #2.

Example 13

Production of chimeric mice from the ES cells retaining human chromosome #2

35

The cells in a frozen stock of the G418 resistant ES cell clone 5-1 that was obtained in Example 12 and which was confirmed to retain human chromosome #2 was thawed, started to culture and injected into 8-cell stage embryos obtained by mating ICR or MCH(ICR) male and female mice (CREA JAPAN, INC.); the injection rate was 10-12 cells per embryo. The embryos were cultured in an ES medium (Example 9) overnight to develop to blastocysts. Two and half days after a foster mother ICR mouse (CREA JAPAN, INC.) was subjected to a pseudopregnant treatment, about ten of the injected embryos were transplanted to each side of the uterus of the foster mother. The results are shown in Table 4.

45

Table 4

Production of chimeric mice from the TT2 cell clone retaining human chromosome #2 (fragments)							
ES cell clone/human chromosome	G418 resistant clone No.	Number of ES cell-injected 8 cell stage embryos	Number of offspring mice	Number of chimeric mice	Contribution to coat color		
TT2/#2 (W23)	5-1	264	51	18	<20%	20-50%	50-80%

As a result of the transplantation of a total of 264 injected embryos, 51 offspring mice were born. Chimerism in the offsprings can be determined by the extent of TT2 cell-derived agouti coat color (dark brown) in the host embryo-

derived albino coat color. Out of the 51 produced offsprings, 18 mice were recognized to have partial agouti coat color, indicating the contribution of the ES cells. The maximum contribution was about 80%.

These results show that the G418 resistant ES cell clone (5-1) retaining a fragment of human chromosome #2 maintains the ability to produce chimera, that is, the ability to differentiate into normal tissues of mouse individual.

5

Example 14

Detection of human antibody heavy chain in sera of the human chromosome #14 transferred chimeric mice

10 The concentrations of human antibody in the sera were determined by enzyme-linked immunosorbent assay (ELISA). The ELISA for human antibody was performed in accordance with the method described in Toyama and Ando, "Monoclonal Antibody Experiment Manual", published by Kodansha, 1987; Andou and Chiba, "Monoclonal Antibody Experiment Procedure Manual", published by Kodansha Scientific, 1991; Ishikawa, "Super High Sensitivity Enzyme Immuno Assay", published by Gakkaisyuppan center, 1993; Ed Harlow and David Lane, "Antibodies A Laboratory Manual", published by Cold Spring Harbor Laboratory, 1988 and A. Doyle and J. B. Griffiths, "Cell & Tissue Culture: Laboratory Procedures", published by John Wiley & Sons Ltd., 1996. In some assays, the condition of reaction were modified, for example, the reaction was performed at 4°C over night. Antibodies to human-immunoglobulin or antigen were diluted to about 0.5-10 µg/ml (100-5000 fold) and ELISA plates were coated with these solutions. PBS supplemented with 5% mouse serum (Sigma, M5905) was used for blocking and dilution of the samples and labeled antibodies. PBS was used for 20-fold dilution of the chimeric mouse sera. After washed, the coated plate was blocked over 1 hour. After plate was washed, sample was added and incubated over a half hour. After washed, Enzyme labeled anti-human immunoglobulin antibodies diluted 100-5000 folds were added to the plates and incubated over 1 hour, the plate was washed and then substrate was added. In some assays, the same procedure was applied except that a biotin-labeled antibody was used. After plate was washed, avidin-enzyme complex was added. After plate was washed, substrate was added. Absorbances were measured with a microplate reader (Bio-tek instrument, EL312e).

15 The chimeric mice (Example 10, K3-1-2, K3-2-2 and K3-2-3) which were 29-35 days old were bled and assayed by ELISA. Anti-human IgM mouse monoclonal antibody (Sigma, I6385) was diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) and absorbed to the 96-well microtiter plates. The serum samples diluted with mouse serum (Sigma, M5905) supplemented PBS were added to the plates. Subsequently, peroxidase-labeled anti-human IgM goat antibody (Tago, 2392) was added and the plates were incubated. After ABTS substrate (Kirkegaard & Perry Laboratories Inc., 506200) was added, enzyme activity was determined by absorbance measurement at 405 nm. Purified human IgM antibody (CAPEL, 6001-1590) and IgG (Sigma, I4506) were used as standards. The standards were diluted stepwise with mouse serum-supplemented PBS. In the determination of human IgG concentration, anti-human IgG goat antibody (Sigma, I3382) was absorbed to the plate and the human IgG was detected with peroxidase-labeled anti-human IgG goat antibody (Sigma, A0170). The results are shown in Table 5. Both human IgM and IgG were detected.

20

25

30

35

Table 5

Concentrations of Human Antibodies in Chimeric Mouse Sera (ELISA)		
Chimeric Mouse	IgG (mg/l)	IgM (mg/l)
K3-1-2	0.37	3.7
K3-2-2	0.33	5.9
K3-2-3	0.51	3.4

40

Two milliliters of human serum albumin (HSA, Sigma, A3782) dissolved in PBS was mixed with adjuvant (MPL+TDM Emulsion, RIBI Immunochem Research Inc.) to prepare an antigen solution at a concentration of 0.25 mg/ml. The chimeric mice retaining human chromosome #14 fragment (Example 10, K3-1-1 and K3-2-1) were immunized with 0.2 ml of the antigen solution 3 times at days 27, 34 and 41 after birth. The chimeric mouse sera were assayed by ELISA. The results are shown in Figs. 13 and 14. The human antibody concentration in the sera of the HSA-immunized chimeric mice was increased after the immunization. In the K3-1-1 mouse, 18 µg/ml of human IgM and 2.6 µg/ml of IgG were detected in the serum at day 17 after the immunization. In the serum of the control ICR mouse, the human antibody titer was not significant.

45

50

55

Example 15

Production of human antibody heavy chain-producing hybridomas from the human chromosome #14 transferred chimeric mouse

5 The spleen was removed from the human albumin-immunized chimeric mouse (K3-1-1, Example 14) at day 44 after birth. The spleen cell was fused with a myeloma cell to produce a hybridoma. The hybridoma was produced using a myeloma cell P3X63Ag8.653 (DAINIPPON PHARMACEUTICAL CO., LTD., 05-565) by the method described in Ando and Chiba, "Monoclonal Antibody Experimental Procedure Manual", published by Kodansha Scientific, 1991. The hybridomas were inoculated into ten 96-well plates and cultured for 1 week. The culture supernatant was analyzed by ELISA. The ELISA procedure was conducted by using antihuman IgM mouse monoclonal antibody (Sigma, I6385) immobilized on ELISA plate in the same manner as in Example 14 to give 6 positive clones. HSA (antigen) was dissolved in 50 mM carbonate-bicarbonate buffer (pH 9.6) at a concentration of 5 µg/ml and the antigen solution was dispensed in 100 µl portions into all the wells of the ELISA plates. After the addition of the supernatant, peroxidase-labeled anti-human IgA+IgG+IgM goat antibodies (Kierkegaard & Perry Laboratories Inc., 04-10-17) were used for detection of HSA-specific human antibody. One positive clone was confirmed in the ten plates. This clone was one of the 6 human IgM positive clones. The clone (H4B7) was further cultured and the culture supernatant was diluted, followed by ELISA analysis using HSA as an antigen with peroxidase-labeled anti-human IgM goat antibody (Tago, 2392) in the same manner as described above. As a result, the absorbance decreased with the increase in the dilution of the culture solution. Serial twofold dilutions of 2 µg/ml human IgM (CAPEL, 6001-1590) showed low absorbance regardless of dilution ratios. This suggests that the antibody produced by hybridoma H4B7 had a specificity to HSA (Fig. 15). In Fig. 15, the dilution of the culture supernatant samples is plotted on the horizontal axis and the absorbance at 405 nm is plotted on the vertical axis.

25 Example 16

Re-marking of the G418 resistance-marked human chromosome #2 fragment with puromycin resistance

30 The A9 cells retaining the G418 resistance-marked human chromosome #2 fragment (W23) (see Example 1, Fig. 1) were cultured in a G418 (800 µg/ml) containing selective medium (10% FBS, DMEM) in a 100 mm plate. Plasmid pPGKPuro (provided by Dr. Peter W. Laird (WHITEHEAD INSTITUTE)) containing puromycin resistance gene was linearized with restriction enzyme SalI (TAKARA SHUZO CO., LTD.) before transfection. The cells were treated with trypsin and suspended in Dulbecco's phosphate buffered saline (PBS) at a concentration of 5×10^6 cells/ml, followed by electroporation using a Gene Pulser (Bio-Rad Laboratories, Inc.) in the presence of 10 µg of DNA in the same manner as in Example 1. A voltage of 1000 V was applied at a capacitance of 25 µF with an Electroporation Cell of 4 mm in length (Example 1) at room temperature. The electroporated cells were inoculated into media in 3-6 plates of 100 mm Ø. After one day, the medium was replaced with a double-selective medium containing 10 µg/ml of puromycin (Sigma, P-7255) and 800 µg/ml of G418. The colonies formed after 2-3 weeks were collected in groups each consisting of about 200 colonies. The cells of each of the three groups were cultured in two or three 25 cm² flasks to form microcells. The 35 mouse A9 cells were cultured in a 25 cm² flask and fused with the microcells by the same procedure as in Example 1. The fused cells were transferred into two 100 mm plates and cultured in the double-selective medium containing G418 and puromycin. One of the three groups gave two double-drug resistant clones. In these clones, it was most likely that puromycin resistance marker had been introduced into human chromosome #2 fragment.

40 Example 17

Duplication of transferred human chromosome in the human chromosome transferred ES cells

45 The ES cell clone retaining the G418 resistance marked human chromosome #14 fragment (E14/#14-36) was cultured in a medium containing G418 at a high concentration to give ES cell clones in which the human chromosome was duplicated ("Biomanual Series 8, Gene Targeting", published by Yodosha, 1995). G418 resistant mouse primary cells (purchased from Lifetech Oriental) were inoculated into a 100 mm plate without treating with mitomycin C and used as feeder cells. The E14/#14-36 cells were inoculated into the 100 mm plate and after half a day, the medium was replaced with a medium containing G418 at a concentration of 16 mg/ml. The medium was replaced every 1-2 days. The G418 50 concentration was changed to 10 mg/ml one week later and the cultivation was continued. Among the colonies formed, 15 were picked up and cultured, followed by FISH analysis of chromosome using human chromosome #14 specific probes (see Example 9). As a result, human chromosome #14 fragment was found to have duplicated in the 8 clones.

Example 18

Preparation of Mouse ES cells retaining both human chromosome #2 partial fragments and human chromosome #14 partial fragments.

5 In a microcell transfer experiment using the double-drug resistant clone PG-1 from Example 16 as a microcell donor cell and a wild-type A9 cell as a recipient cell, it was confirmed that the human chromosome #2 partial fragment retained in PG-1 was marked with a puromycin resistance gene. The preparation of microcells and the fusion with the A9 cells was carried out by the same methods as in Example 1. As a result, 10 days after the microcell fusion, a total
10 of fifty nine G418 resistant colonies appeared. After the medium for these colonies was changed to one containing 8 µg/ml puromycin, the colonies were cultured for 3 days to give 45 viable colonies (76%). In many cases of microcell fusion, only one or few chromosomes are transferred into a recipient cell. Hence, cotransfer of both the resistance genes at a high frequency shows that the G418 resistance-labeled chromosome #2 partial fragment retained in the PG1 clone was also marked with the puromycin resistance gene. In addition, for the detection of the respective marker
15 genes on the human chromosome #2 partial fragment, FISH analysis was conducted by using pSTneoB (see Example 1) as a probe in the case of the A9/#2 W23 clone having only G418 resistance (see Example 16) and by using pPGKPuro (see Example 16) as a probe in the case of the PG1 clone in accordance with the method described in Matsubara et al., "FISH Experiment Protocol", published by Shujunsha, 1994. As a result, in the case of the A9/#2 W23 clone,
20 one signal was observed in each of the sister chromatids of the human chromosome #2 partial fragment observed in Example 12 (2 signals in total). This indicated the insertion of pSTneoB into the human chromosome #2 partial fragment at one site. In the case of the PG1 clone, a total of 4 signals were observed on a chromosome fragment of the same size as in A9/#2 W23. Since pSTneoB and pPGKPuro had identical sequences in their vector portions, the pSTneoB could be detected by the pPGKPuro probe. Hence, it is believed that out of the four signals observed in the PG1 clone, two were from the pSTneoB and the other two were from the pPGKPuro. These results show that the human
25 chromosome #2 partial fragment retained in the PG1 was marked with both the G418 and puromycin resistances.

The PG1 cell clone was used as a chromosome donor cell to prepare a mouse ES cell retaining both a human chromosome #2 partial fragment and a human chromosome #14 partial fragment. The G418 resistant TT2 cell clone 1-4 already retaining the human chromosome #14 partial fragment (see Example 9) was used as a chromosome recipient cell. The microcell fusion and the selection of puromycin resistant cells were carried out by the same methods as in the
30 selection of the G418 resistant clones in Example 9 except that the concentration of puromycin was 0.75 µg/ml. The frequency of the appearance of the resulting puromycin resistant clones was 3-7 per 10⁷ of 1-4 cells. The presence of G418 resistance in these puromycin resistant clones was confirmed from the fact that they were grown in the presence of 300 µg/ml of G418. The double-drug resistant clones were stored frozen and genomic DNA was prepared by the same methods as in Example 2. The retention of the human chromosome #2 partial fragment and human chromosome
35 #14 partial fragment was confirmed by the method described in (1) in the case of double-drug resistant clones PG5, PG15 and PG16 and by the method described in (2) in the case of the clone PG15.

(1) PCR analysis

40 Genomic DNAs of the double-drug resistant clones were used as templates in the PCR amplifications. Among the markers on human chromosomes #2 and #14 (Genetic Maps, supra), the primers whose presence in the A9/#2 W23 clone was confirmed in Example 12 and those whose presence in the TT2/#14 1-4 clone was confirmed in Example 9 were used. All the primers gave expected amplification products in all the three clones.

(2) Fluorescence in situ hybridization (FISH)

FISH analysis was conducted by using FITC-labeled human total DNA as a probe in the same manner as in Example 11. As a result, in almost all of the metaphase spreads, two (large and small) human chromosome fragments were detected. The large fragment had the same size as that of the partial fragment detected by using the human chromosome #14 specific probes in the case of the TT2/#14 1-4 clone in Example 9 and the small fragment had the same size as that of the partial fragment detected by using the human chromosome #2 specific probes in the case of the TT2/#2 5-1 in Example 12. The results are shown in Fig. 16. In Fig. 16, the less bright chromosome was derived from the mouse. The two (large and small) chromosome fragments of high brightness due to FITC fluorescence as shown by arrows were derived from the human, which are believed to correspond to the human chromosome #14 and #2 partial fragments.
55

These results show that the obtained double-drug resistant ES clones retained both the human chromosome #2 partial fragment and the human chromosome #14 partial fragment.

Example 19

Production of chimeric mice from the mouse ES cell clones retaining both human chromosome #2 partial fragments and human chromosome #14 partial fragments

The cells in frozen stocks of the G418 and puromycin double-resistant TT2 cell clones PG5, PG15 and PG16 from Example 18 which were confirmed to retain human chromosome #2 partial fragments and human chromosome #14 partial fragments were thawed, started to culture and injected into 8-cell stage embryos obtained by mating ICR or MCH(ICR) male and female mice (CREA JAPAN, INC.); the injection rate was 10-12 cells per embryo. The embryos were cultured in a medium for ES cells (see Example 9) overnight to develop to blastocysts. Two and a half day after a foster mother ICR mouse was subjected to a pseudopregnant treatment, about ten of the injected embryos were transplanted to each side of the uterus of the foster mother. The results are shown in Table 6.

15

Table 6

20

Production of chimeric mice from the mouse ES cell clones retaining both human chromosome #2 partial fragments and human chromosome #14 partial fragments					
ES cell clone/human chromosome	Double-drug resistant clone No.	Number of ES cell-injected 8-cell stage embryos	Number of off-spring mice	Number of chimeric mice	Contribution to coat color
					<10% 10-50% 50%<
TT2/#14+#2	PG5	160	26	8	7 1 -
	PG15	168	15	3	1 2 -
	PG16	223	32	12	3 6 3

30

As a result of the transplantation of a total of 551 injected embryos, 73 offspring mice were born. Chimerism in the offsprings can be determined by the extent of TT2 cell-derived agouti coat color (dark brown) in the host embryo-derived albino coat color. Out of the 73 produced offsprings, 23 mice were recognized to have a partial agouti coat color, indicating the contribution of the ES cells.

35

These results show that the ES cell clones PG5, PG15 and PG16 retaining human chromosome #2 partial fragments and human chromosome #14 partial fragments maintain the ability to produce chimera, that is, the ability to differentiate into normal tissues of mouse.

40

Example 20
Detection of human antibody in sera of the chimeric mice derived from the ES cells retaining both human chromosome #2 partial fragments and human chromosome #14 partial fragments

45

The two KPG-15 (9 weeks old; derived from the PG-5 clone, 10% chimerism) and KPG-18 (5 weeks old; derived from the PG-5 clone, 10% chimerism) chimeric mice from Example 19 were immunized with 0.2 ml of a solution of human serum albumin (HSA, Sigma, A3782) and adjuvant (MPL+TDM Emulsion, RIBI Immunochem Research Inc.) at a HSA concentration of 0.25 mg/ml. The chimeric mice were bled just before the immunization and 8 days after that and the concentrations of human antibody μ and κ chains in the sera were determined by ELISA (see Example 14). Ninety six-well microtiter plates were coated with antihuman antibody κ chain goat antibody (VECTOR LABORATORIES INC., AI-3060) diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) and then a serum sample diluted with mouse serum (Sigma, M5905)-containing PBS was added. Subsequently, biotin-labeled anti-human antibody κ chain goat antibody (VECTOR LABORATORIES INC., BA-3060) was added to the plates and incubated. A complex of biotinylated horse-radish peroxidase and avidin DH (VECTOR LABORATORIES, INC., Vectastain ABC Kit, PK4000) was added and incubated. After 3,3',5,5'-tetramethylbenzidine (TMBZ, Sumitomo Bakelite, ML-1120T) was added as a peroxidase substrate, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgG antibody having κ chain (Sigma, I-3889) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS. In the case of μ chain, 96-well microtiter plates were coated with anti-human antibody μ chain mouse monoclonal antibody (Sigma, I-6385) diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) and then a serum sample was added. Subsequently, peroxidase-labeled anti-human antibody μ chain mouse antibody (The Binding Site Lim-

ited, MP008) was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgM antibody having μ chain (CAP-PEL, 6001-1590) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS. As a result, both the human antibody μ and κ chains were detected in both individuals. The concentrations of these human antibodies in the sera increased after the immunization (Tables 7 and 8).

Table 7

Concentrations of Human Antibodies in Chimeric Mouse KPG15 (ELISA)		
	IgM (mg/l)	Ig κ (mg/l)
Before Immunization	0.19	1.6
8 Days After Immunization	0.75	1.7

Table 8

Concentrations of Human Antibodies in Chimeric Mouse KPG18 (ELISA)		
	IgM (mg/l)	Ig κ (mg/l)
Before Immunization	0.29	0.57
8 Days After Immunization	3.4	0.87

These results show that human antibody heavy and light chain genes can function in the chimeric mice derived from the ES cells retaining both human chromosome #2 partial fragments and human chromosome #14 partial fragments.

Example 21

Detection of anti-HSA human antibody γ chain in sera of the human chromosome #14 fragments transferred chimeric mice

The chimeric mice retaining human chromosome #14 fragments which were produced by the same method as in Example 10 (K9 and K11: both were derived from the TT2 cell clone 3-2, with chimerisms of 50% and 30%, respectively) were immunized with HSA either 4 times at days 79, 93, 107 and 133 after birth (K9) or 3 times at days 74, 88 and 111 after birth (K11) by the same method as in Example 20. Antibodies including human γ chain against human serum albumin in the sera of the chimeric mice were detected by ELISA. Ninety six-well microtiter plates were coated with HSA (Sigma, A 3782) diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) and then a sample diluted with PBS was added. Subsequently, peroxidase-labeled anti-human IgG mouse antibody (Pharmingen, 08007E) was added to the plates and incubated. After Ophenylenediamine (OPD, Sumitomo Bakelite, ML-1130O) was added as a peroxidase substrate, enzyme activity was determined by absorbance measurement at 490 nm. The titer of the anti-HSA human IgG in the sera of the chimeric mice immunized with HSA increased after the immunization. On the other hand, control ICR mouse gave a background level of the anti-HSA human IgG titer after the immunization with HSA. The results are shown in Fig. 17. In Fig. 17, the number of days after the first immunization of the chimeric mice with HSA is plotted on the horizontal axis and the absorbance at 490 nm is plotted on the vertical axis. These results show that the antibody titer of the antigen specific human IgG was increased by stimulation with the HSA antigen in the chimeric mice retaining human chromosome #14 fragments.

Example 22

Detection of human antibody λ chain in a serum of the human chromosome #22 fragment transferred chimeric mouse

The chimeric mouse K22-7 from Example 3 (9 weeks old; 10% chimerism) was bled and human antibody λ chain

in the serum was detected by ELISA (see Example 14). Ninety six-well microtiter plates were coated with anti-human antibody λ chain goat antibody (VECTOR LABORATORIES INC., AI-3070) diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) and then a serum sample was added. Subsequently, biotin-labeled anti-human antibody λ chain goat antibody (VECTOR LABORATORIES INC., BA-3070) was added to the plates and incubated. A complex of biotinylated horseradish peroxidase and avidin DH (VECTOR LABORATORIES, INC., Vectastain ABC Kit) was added and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added as a peroxidase substrate, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgG antibody having λ chain (Sigma, I-4014) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS. As a result, human antibody λ chain was detected in the chimeric mouse at a concentration corresponding to 180 ng/ml of human IgG. These results show that human antibody λ chain gene can function in the chimeric mouse retaining a human chromosome #22 fragment.

Example 23

15 Detection of human antibody κ chain in sera of the human chromosome #2 fragment transferred chimeric mice

The chimeric mouse K2-8 from Example 13 (5 weeks old; 70% chimerism) and the chimeric mice K2-3, K2-4 and K2-12 from Example 13 (9 weeks old; chimerisms was 50%, 20% and 80%, respectively) were bled and human antibody κ chain in the sera was detected by ELISA (see Example 14). Ninety six-well microtiter plates were coated with antihuman antibody κ chain goat antibody (VECTOR LABORATORIES INC., AI-3060) diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) and then a serum sample was added. Subsequently, biotin-labeled anti-human antibody κ chain goat antibody (VECTOR LABORATORIES INC., BA-3060) was added to the plates and incubated. A complex of biotinylated horseradish peroxidase and avidin DH (VECTOR LABORATORIES, INC., Vectastain ABC Kit) was added and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgG antibody having κ chain (Sigma, I-3889) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS. The results are shown in Table 9.

Table 9

Concentration of Human Anti-body κ Chain in Chimeric Mouse (ELISA)	
Chimeric Mouse	Ig κ (mg/l)
K2-3	124
K2-4	85
K2-8	25
K2-12	56

The chimeric mice K2-3 and K2-4 retaining human chromosome #2 fragments from Example 13 were immunized with HSA, 3 times at days 66, 80 and 102 after birth by the same method as in Example 20. The chimeric mouse K2-12 was immunized with HSA, 4 times at days 63, 77, 91 and 116 after birth by the same method as in Example 20. Human antibody κ chain against HSA in the sera of the chimeric mice was detected by ELISA (see Example 14). Ninety six-well microtiter plates were coated with HSA (Sigma, A 3782) diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) and then a sample was added. Subsequently, biotin-labeled anti-human antibody κ chain goat antibody (VECTOR LABORATORIES, INC., BA-3060) was added to the plates and incubated. A complex of biotinylated horseradish peroxidase and avidin DH (VECTOR LABORATORIES, INC., Vectastain ABC Kit) was added and incubated. After OPD (Sumitomo Bakelite, ML-1130O) was added as a peroxidase substrate, enzyme activity was determined by absorbance measurement at 490 nm. The titer of the anti-HSA human κ chain in the sera of the chimeric mice immunized with HSA increased after the immunization. On the other hand, control ICR mouse gave a background level of the anti-HSA human κ chain titer after the immunization. The results are shown in Fig. 18. In Fig. 18, the number of days after the first immunization of the chimeric mice with HSA is plotted on the horizontal axis and the absorbance at 490 nm is plotted on the vertical axis. These results show that human antibody κ chain gene can function in the chimeric mice retaining human chromosome #2 fragments and that the antibody titer of the antigen specific human Ig κ was increased by stimulation with the HSA antigen in the chimeric mice.

Example 24

Preparation of human antibody heavy chain (μ chain or γ chain)-producing hybridomas from the human chromosome #14 transferred chimeric mouse

5 The spleen was removed from the HSA-immunized chimeric mouse K9 (see Example 21) at day 136 after birth. A spleen cell was fused with a myeloma cell to produce a hybridoma. The hybridoma was produced using a myeloma cell Sp-2/0-Ag14 (Dainippon Pharmaceutical Co., Ltd., 05-554) by the method described in Toyama and Ando, "Monoclonal Antibody Experiment Procedure Manual", published by Kodansha Scientific, 1991. The cells were inoculated into a
10 medium containing 10% ORIGEN Hybridoma Cloning Factor (HCF, Bokusui Brown) in eight 96-well plates and G418 was added after 3 days at a concentration of 1 mg/ml, followed by cultivation for 1-3 weeks. The culture supernatant was analyzed by ELISA. Ninety six-well microtiter plates were coated with anti-human μ chain mouse monoclonal antibody (Sigma, I-6385) diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) and a sample diluted with PBS was added. Subsequently, peroxidase-labeled anti-human μ chain mouse antibody (The Binding Site LIMITED, MP008) was
15 added to the plates and incubated. 2,2'-Azino-di-(3-ethyl-benzothiazoline-6-sulfonate) diammonium salt (ABTS, Kirkegaard & Perry Laboratories Inc., 04-10-17) was used as a substrate to detect seven positive clones. In the detection of γ chain-producing clones, 96-well microtiter plates were coated with anti-human γ chain mouse monoclonal antibody (Sigma, I-6260) and a sample diluted with PBS was added. Subsequently, peroxidase-labeled anti-human γ chain mouse antibody (Pharmingen, 08007E) was added to the plates and incubated. ABTS (Kirkegaard & Perry Laboratories Inc., 04-10-17) was used as a substrate and two human antibody γ chain-positive clones were obtained.
20

Example 25

Preparation of human antibody light chain-producing hybridomas from the human chromosome #2 transferred chimeric mouse

25 The spleen was removed from the HSA-immunized chimeric mouse K2-3 (see Example 23) at day 105 after birth. A spleen cell was fused with a myeloma cell to produce a hybridoma. The hybridoma was produced using a myeloma cell P3X63Ag8.653 (Dainippon Pharmaceutical Co., Ltd., 05-565) by the method described in Ando and Chiba, "Monoclonal Antibody Experiment Procedure Manual", published by Kodansha Scientific, 1991. The cells were inoculated into a medium containing 10% HCF (Bokusui Brown) in ten 96-well plates and G418 was added after 3 days at a concentration of 1 mg/ml, followed by cultivation for 1-3 weeks. The culture supernatant was assayed by ELISA. The ELISA analysis was conducted by the same method as in Example 23 and two human antibody κ chain-positive clones were obtained.
35

Example 26

Re-marking of the G418 resistance-marked human chromosome #22 with puromycin resistance

40 The A9 cells retaining the G418 resistance-marked human chromosome #22 (A9/#22 γ 2) from Example 1 were re-marked with puromycin resistance by the same method as in Example 16. About 200 colonies of double-drug resistant clones obtained by electroporation of the γ 2 cells with pPGXPuro were collected as one group and three such groups (P1, P2 and P3) were used as donor cells to perform microcell transfer into wild-type mouse A9 cells. As a result, 6, 1 and 3 of double-drug resistant clones were obtained from the groups P1, P2 and P3, respectively. The clone 6-1 from
45 group P3 was used as a microcell donor cell and a wild-type A9 cell as a recipient cell to perform a microcell transfer experiment (see Example 18). As a result, the human chromosome #22 was confirmed to have been further marked with a puromycin resistance gene. The preparation of microcells and the fusion with A9 cells were conducted by the same methods as in Example 1. As a result, twenty eight G418 resistant colonies appeared 11 days after the microcell transfer. After the medium for these colonies was changed to one containing 8 μ g/ml puromycin, these colonies were
50 cultured for 3 days to give 21 (75%) viable colonies. In many cases of microcell fusion, only one or few chromosomes are transferred into a recipient cell. Hence, cotransfer of both the resistance genes at a high frequency shows that the G418 resistance-labeled chromosome #22 retained in the 6-1 clone was marked with the puromycin resistance gene.

Example 27

55 Preparation and Sequencing of cDNA of a human antibody heavy chain variable region from the human antibody heavy chain-producing hybridoma

Among the human antibody heavy chain (IgM)-producing hybridomas obtained in Example 15, H4B7 (HSA-spe-

cific) and H8F9 (non-specific) hybridomas were selected. Total RNAs were obtained from these hybridomas using ISOGEN (Nippon Gene). The synthesis of cDNA from 5 µg each of the total RNAs was conducted with a Ready-To-Go T-primed 1st strand Kit (Pharmacia Co.). Using the resulting cDNA and the following primers prepared with reference to Larrick et al., BIO/TECHNOLOGY, 7, 934-, 1989; Word et al., Int. Immunol., 1, 296-, 1989, PCR was performed to amplify a human antibody heavy chain variable region.

5 CM1 (human IgM constant region) : 5'-TTGTATTCAGGAGAAAGTG (SEQ ID NO: 45)

CM2 (ditto) : 5'-GGAGACGAGGGGAAAGGG (SEQ ID NO:46)

HS1 (human heavy chain variable region) : 5'-ATGGACTGGACCTGGAGG(AG) TC(CT)TCT(GT)C (SEQ ID NO:47) (a mixture of 8 sequences)

10 HS2 (ditto) : 5'-ATGGAG(CT)TTGGGCTGA(GC)CTGG(GC)TTT(CT)T (SEQ ID NO:48) (a mixture of 16 sequences)

HS3 (ditto) : 5'-ATG(AG)A(AC)(AC)(AT)ACT(GT)TG(GT)(AT)(GCT)C(AT)(CT) (GC)CT(CT)CTG (SEQ ID NO:49) (a mixture of 6144 sequences)

In both cases of the H4B7 and H8F9 hybridomas, the first run of PCR was performed by using three kinds of primer combinations of HS1xCM1, HS2xCM1 and HS3xCM1 in 40 cycles at 94 °C for 1 minute, 50 °C for 2 minutes and 72 °C

15 for 3 minutes with a Thermal Cycler 140 (Perkin-Elmer Corp.). The PCR products were amplified again under the same temperature conditions in 30 cycles using HS1xCM2, HS2xCM2 and HS3xCM2 primers, respectively. The amplification products were electrophoresed on a 1.5% agarose gel and detected by staining with ethidium bromide. As a result, an amplification product of about 490 bp was detected with the HS3xCM2 primer in the case of the H4B7 hybridoma. In the case of the H8F9 hybridoma, a slight band was detected at the same site with the HS3xCM2 primer. The band in

20 the case of H8F9 was amplified again with the HS3xCM2 primer in 30 cycles under the same temperature conditions as above. As a result, the amplification product was detected as a very intensive signal. These PCR products were cloned into a pBlueScriptII SK+ (Stratagene Ltd.) at a SmaI site in accordance with the method described in Ishida et al., "Gene Expression Experiment Manual", published by Kodansha Scientific, 1995. Among the amplification product-

25 inserted plasmids, plasmids #2, #3, #4(H4B7), #11, #13 and #14 (H8F9) were selected and the nucleotide sequences of the amplification products were determined with a Fluorescence Autosequencer (Applied Biosystems Inc.). As a result of the comparison of the obtained nucleotide sequences or deduced amino acid sequences with those of known human antibody VH region (Marks et al., Eur. J. Immunol. 21, 985-, 1991) and JH region (Ravetch et al., Cell, 27, 583-, 1981), it was revealed that both the H4B7 and H8F9 hybridomas contained a combination of genes for VH4 family and JH2. These results show that the chimeric mouse retaining human chromosome #14 partial fragment produced a complete functional human antibody heavy chain protein.

Example 28

Preparation and Sequencing of cDNA of human antibody κ chain from the spleen of the human antibody κ chain-expressing chimeric mouse

In the same manner as in Example 5, cDNA was prepared from the spleen of the chimeric mouse K2-8 from Example 13 which was confirmed to express human antibody κ chain in Example 23. Using the resulting cDNA and the following primers prepared with reference to Larrick et al., BIO/TECHNOLOGY, 7, 934-, 1989; Whitehurst et al., Nucleic Acids Res., 20, 4929-, 1992, PCR was performed to amplify human antibody κ chain variable region. cDNA from the liver of the chimeric mouse K2-8 and cDNA from the spleen of the chimeric mouse K3-2-2 derived from the TT2/#14 3-2 clone (see Example 10) were used as negative controls.

KC2 (human Ig κ chain constant region) : 5'-CAGAGGCAGTCCAGATTTC (SEQ ID NO:50)

KC3 (ditto) : 5'-TGGGATAGAAGTTATTTCAGC (SEQ ID NO:51)

45 KVMIX (human Ig κ chain variable region) : 5'-ATGGACATG(AG)(AG)(AGT)(CT)CC(ACT)(ACG)G(CT)(GT)CA(CG)CTT (SEQ ID NO:52) (a mixture of 3456 sequences)

PCR was performed by using primer combinations of KVMIXxKC2 and KVMIXxKC3 in 40 cycles at 94 °C for 15 seconds, 55 °C for 15 seconds and 72°C for 20 seconds with a Thermal Cycler 9600 (Perkin-Elmer Corp.). The amplification products were electrophoresed on a 1.5% agarose gel and detected by staining with ethidium bromide. As a result, expected amplification products of about 420 bp (KC2) and about 450 bp (KC3) were detected. In the case of the two negative controls, no specific amplification product was detected. These amplification products were cloned into a pBlueScriptII SK+ (Stratagene Ltd.) at a SmaI or EcoRI site in accordance with the method described in Ishida et al., "Gene Expression Experiment Manual", published by Kodansha Scientific, 1995. Among the amplification product-inserted plasmids, VK-#1 clone derived from the KVMIXxKC2 primers was selected and the nucleotide sequence of the amplification product was determined with a Fluorescence Autosequencer (Applied Biosystems Inc.). Since the obtained nucleotide sequence did not contain a termination codon at any site between an initiation codon and a con-

* () means that any one of the bases therein should be selected.

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stant region of human Ig κ chain, the cloned amplification products are believed to encode a variable region of functional human Ig κ chain. As a result of the comparison of the obtained nucleotide sequences with those of known human antibody V κ region (Klein et al., Eur. J. Immunol. 23, 3248, 1993) and J κ region (Whitehurst et al., supra), it was revealed that the VK-#1 clone contained a combination of genes for V κ 3 family and J κ 4. These results show that the chimeric mouse retaining human chromosome #2 partial fragment produced a complete functional human antibody κ chain protein.

Example 29

10 Detection and quantitation of human antibody γ chain subclasses and μ chain in sera of the chimeric mice retaining human chromosome #14 fragment

The chimeric mice K15A and K16A from Example 10 (derived from the 1-4 clone, with chimerism of 70% and 50%, respectively) of 11 weeks after birth were bled and human antibody γ chain subclasses and μ chain in the sera were detected by the same ELISA method as in Example 14.

Quantitative Determination of human IgG1

Ninety six-well microtiter plates were coated with anti-human IgG antibody (Sigma, I-6260) diluted with PBS. A serum sample was added. Subsequently, peroxidase-labeled anti-human IgG1 antibody (Pharmingen, 08027E) was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgG1 antibody (Sigma, I-3889) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS.

25 **Quantitative Determination of human IgG2**

Ninety six-well microtiter plates were coated with anti-human IgG2 antibody (Sigma, I-9513) diluted with PBS. A serum sample was added. Subsequently, peroxidase-labeled anti-human IgG antibody (Sigma, A-0170) was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgG2 antibody (Sigma, I-4139) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS.

Quantitative Determination of human IgG3

35 Anti-human IgG3 antibody (Sigma, I-7260) was diluted with 100 mM glycine-HCl buffer (pH 2.5) and incubated for 5 minutes at room temperature, followed by 10-fold dilution with 100 mM phosphate buffer (pH 7.0). Ninety six-well microtiter plates were coated with the antihuman IgG3 antibody solution. A serum sample was added. Subsequently, peroxidase-labeled anti-human IgG antibody (Pharmingen, 08007E) was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgG3 antibody (Sigma, I-4389) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS.

Quantitative Determination of human IgG4

45 Anti-human IgG4 antibody (Sigma, I-7635) was diluted with 100 mM glycine-HCl buffer (pH 2.5) and incubated for 5 minutes at room temperature, followed by 10-fold dilution with 100 mM phosphate buffer (pH 7.0). Ninety six-well microtiter plates were coated with the anti-human IgG3 antibody solution. A serum sample was added. Subsequently, peroxidase-labeled anti-human IgG antibody (Pharmingen, 08007E) was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgG4 antibody (Sigma, I-4639) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS.

Quantitative Determination of human IgM

55 Ninety six-well microtiter plates were coated with anti-human μ chain mouse monoclonal antibody (Sigma, I-6385) diluted with PBS. A serum sample was added. Subsequently, peroxidase-labeled anti-human μ chain mouse antibody (The Binding Site Limited, MP008) diluted with mouse serum (Sigma, M5905)-supplemented PBS was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added as a peroxidase substrate, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgM having μ chain (CAPPEL, 6001-1590)

was used as standard. The standard was diluted stepwise with mouse serum (Sigma, M5905)-supplemented PBS.

The results are shown in Table 10. All the subclasses IgG1, IgG2, IgG3 and IgG4, and IgM were detected in the two chimeric mice K15A and K16A.

5

Table 10

Concentrations of Human antibody IgG Subclasses and IgM in the Chimeric Mice (ELISA)					
Chimeric mouse	IgG1	IgG2	IgG3	IgG4	IgM (mg/l)
K15A	2.25	1.96	0.17	0.43	7.09
K16A	0.30	0.69	0.10	0.07	0.87

15

Example 30

Preparation of mouse ES cell clones (TT2) retaining human chromosome #22

20

The cell clone 6-1 (A9/#22, G418 and puromycin resistant) from Example 26 was used as a chromosome donor cell for the preparation of mouse ES cell (TT2) retaining human chromosome #22. A wild-type TT2 cell line (see Example 9) was used as a chromosome recipient cell. The microcell fusion and the selection of puromycin resistant clones were conducted by the same procedures as in the selection of G418 resistant clones in Example 9 except that the concentration of puromycin was 0.75 µg/ml. The frequency of the appearance of the puromycin resistant clones was 1-2 per 10⁷ of TT2 cells. The puromycin resistant clones were stored frozen and genomic DNA was prepared by the same methods as in Example 2. The retention of human chromosome #22 in the puromycin resistant clone PG22-1 was confirmed by the methods described in (1) and (2) below.

30 (1) PCR analysis

Genomic DNA of the puromycin resistant clone was used as a template in PCR amplification. Among the genes on human chromosome #22 (Genetic Maps, supra), ten primers whose presence in the A9/#22 clone was confirmed in Example 2 were used in the PCR amplification. All the markers which existed in the A9/#22 clone (see Example 2) were detected.

35 (2) Southern blot analysis

In accordance with the same method as described in Example 2 using human L1 sequence as a probe, Southern blot analysis was conducted with genomic DNAs obtained from wild-type TT2 (negative control), the chromosome donor cell 6-1 and the puromycin resistant TT2 cell clone PG22-1. The results are shown in Fig. 19. In Fig. 19, the molecular weights of DNAs are shown at the left side. The band pattern of the PG22-1 clone was equivalent to that of the 6-1 cell and the signal intensities were the same. Hence, it was confirmed that chromosome #22 in the 6-1 cell had been transferred certainly into the PG22-1 clone.

40 These experiments demonstrate that the puromycin resistant TT2 cell clone PG22-1 retained the whole or the most part of human chromosome #22.

Example 31

Production of chimeric mice from the mouse ES cells (TT2) retaining human chromosome #22

The cells in a frozen stock of the puromycin resistant TT2 cell clone PG22-1 from Example 30 which was confirmed to retain human chromosome #22 were thawed, started to culture and injected into 8-cell stage embryos obtained by mating ICR or MCH(ICR) male and female mice (CREA JAPAN, INC.); the injection rate was 10-12 cells per embryo. The embryos were cultured in a medium for ES cells (see Example 9) overnight to develop to blastocysts. Two and a half day after a foster mother ICR mouse was subjected to a pseudopregnant treatment, about ten of the injected embryos were transplanted to each side of the uterus of the foster mother. The results are shown in Table 11.

Table 11

Production of chimeric mice from the TT2 cell clone retaining human chromosome #22							
	ES cell clone/human chromosome	Puromycin resistant clone No.	Number of ES cell-injected 8-cell stage embryos	Number of off-spring mice	Number chimeric mice	Contribution to coat color	
10						<20%	20-50%
TT2/#22	PG22-1	266	36	8	4	1	3

15 As a result of the transplantation of a total of 266 injected embryos, 36 offspring mice were born. Chimerism in the offsprings can be determined by the extent of TT2 cell-derived agouti coat color (dark brown) in the host embryo-derived albino coat color. Out of the 36 produced offsprings, 8 mice were recognized to have a partial agouti coat color, indicating the contribution of the ES cells.

20 These results show that the ES cell clone (derived from TT2, PG22-1) retaining human chromosome #22 maintain the ability to produce chimera, that is, the ability to differentiate into normal tissues of mouse.

Example 32

Detection and quantitation of human antibody λ chain in sera of the chimeric mice retaining human chromosome #22

25 The concentration of human antibody λ in the sera of the chimeric mice KPG22-1, 2 and 3 from Example 31 was determined by ELISA in accordance with the same procedure as in Example 14. The chimeric mice of 2 months after birth were bled and human antibody λ chain in the sera was detected by ELISA. Ninety six-well microtiter plates were coated with anti-human immunoglobulin λ chain antibody (VECTOR LABORATORIES INC., IA-3070) diluted with PBS 30 and then a serum sample was added. Subsequently, biotin-labeled anti-human immunoglobulin λ chain antibody (VECTOR LABORATORIES INC., BA-3070) was added to the plates and incubated. A complex of biotinylated horseradish peroxidase and avidin DH (VECTOR LABORATORIES, INC., Vectastain ABC Kit) was added and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgM antibody having λ chain (Dainippon Pharmaceutical Co., Ltd., U13200) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS. The results are shown in Table 12. These results show that human antibody λ chain gene can function in the chimeric mice retaining human chromosome #22.

Table 12

Concentration of Human Antibody λ Chain in Chimeric Mice (ELISA)		
Chimeric Mouse	%Chimerism	Ig λ (mg/l)
KPG22-1	50	12
KPG22-2	50	18
KPG22-3	20	24

50

Example 33

55 Detection of anti-human HSA human antibody λ chain in a serum of the human chromosome #22 transferred chimeric mouse

The chimeric mouse KPG22-3 from Example 31 was immunized with HSA, 3 times at days 79, 94 and 110 after birth by the same method as in Example 20. Human antibody λ chain in the serum of the chimeric mouse was detected

by ELISA in accordance with the same procedure as in Example 14. Ninety six-well microtiter plates were coated with HSA (Sigma, A 3782) diluted with 50 mM carbonate-bicarbonate buffer (pH 9.6) to a concentration of 5 µg/ml and a serum sample was added. Biotinylated anti-human Ig λ antibody (VECTOR LABORATORIES INC., BA-3070) was added. Subsequently, a complex of biotinylated-horseradish peroxidase and avidin DH (VECTOR LABORATORIES, INC., Vectastain ABC Kit) was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added, enzyme activity was determined by absorbance measurement at 450 nm. The titer of the anti-HSA human λ chain in the serum of the chimeric mouse increased after the immunization. On the other hand, control ICR mouse gave a background level of the anti-HSA human λ chain titer after the immunization with HSA. The results are shown in Fig. 20. In Fig. 20, the number of days after the first immunization of the chimeric mouse with HSA is plotted on the horizontal axis and the absorbance at 450 nm is plotted on the vertical axis. These results show that human antibody λ chain gene can function in the chimeric mouse retaining human chromosome #22 and that the antibody titer of the antigen specific human Igλ was increased by stimulation with the HSA antigen.

Example 34

15

Preparation of human antibody light chain-producing hybridomas from the human chromosome #22 transferred chimeric mouse

The spleen was removed from the mouse KPG22-3 (see Example 33) at day 113 after birth by the same method as in Example 25. A spleen cell was fused with a myeloma cell to produce a hybridoma. The hybridoma was produced using a myeloma cell SP-2/0-Ag14 (Dainippon Pharmaceutical Co., Ltd., 05-554) by the method described in Ando and Chiba, "Monoclonal Antibody Experiment Manual", published by Kodansha Scientific, 1991. The cells were inoculated into a medium containing 10% HCF (Air Brown) in five 96-well plates and cultured for 1-3 weeks. The supernatant of the culture solution in colony-positive wells was analyzed by ELISA. The ELISA analysis was conducted by the same method as in Example 33 and four human antibody λ chain-positive clones were obtained.

Example 35

30

Preparation of mouse ES cell clones retaining both a human chromosome #22 partial fragment and a human chromosome #14 partial fragment

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The 6-1 cell clone from Example 26 (A9/#22, G418 and puromycin resistant) was used as a chromosome donor cell for the preparation of mouse ES cells retaining both a human chromosome #22 partial fragment and a human chromosome #14 partial fragment. The G418 resistant TT2 cell clone 1-4 retaining a human chromosome #14 partial fragment from Example 9 was used as a chromosome recipient cell. The experiment of microcell fusion and the selection of puromycin resistant cells were carried out by the same methods as in the selection of the G418 resistant clones in Example 9 except that the concentration of puromycin was 0.75 µg/ml. As a result, the frequency of the appearance of the puromycin resistant clones was 1-2 per 10⁷ of 1-4 cells. The retention of G418 resistance in the puromycin resistant clones was confirmed from the fact that these clones were grown in the presence of 300 µg/ml G418. The double-drug resistant clones were stored frozen and genomic DNAs were prepared by the same methods as in Example 2. The retention of human chromosome #22 and a human chromosome #14 partial fragment in the double-drug resistant clone PG22-5 was confirmed by PCR analysis. With genomic DNA of the double-drug resistant clone used as a template, PCR amplification was conducted using primers whose presence on chromosome #22 was confirmed in Example 2 (A9/#22) and primers whose presence on chromosome #14 was confirmed in Example 9 (TT2/#14 1-4); as a result, three markers (D22S275, D22S315 and Igλ) of the ten markers on chromosome #22 and all of the markers on chromosome #14 in the TT2/#14 1-4 clone were detected.

40

These experiments demonstrate that the obtained double-drug resistant TT2 cell clone retained both a human chromosome #22 partial fragment and a human chromosome #14 partial fragment.

45

Example 36

Production of the chimeric mouse from the mouse ES cell clone retaining both a human chromosome #22 partial fragment and a human chromosome #14 partial fragment

55

The cells in a frozen stock of the G418 and puromycin double-resistant TT2 cell clone PG22-5 from Example 35 which was confirmed to retain a human chromosome #22 partial fragment and a human chromosome #14 partial fragment were thawed, started to culture and injected into 8-cell stage embryos obtained by mating ICR or MCH(ICR) male and female mice (CREA JAPAN, INC.); the injection rate was 10-12 cells per embryo. The embryos were cultured in a medium for ES cells (see Example 9) overnight to develop to blastocysts. Two and a half day after a foster mother ICR

mouse was subjected to a pseudopregnant treatment, about ten of the injected embryos were transplanted to each side of the uterus of the foster mother. The results are shown in Table 13.

Table 13

Production of the chimeric mouse from the mouse ES cell clone retaining both a human chromosome #22 partial fragment and a human chromosome #14 partial fragment								
	ES cell clone/human chromosome	Double-drug resistant clone No.	Number of ES cell-injected 8-cell stage embryos	Number of off-spring mice	Number of chimeric mice	Contribution to coat color		
						<20%	20-50%	50-80%
15	TT2/#22+/#14	PG22-5	302	16	5	3	2	0

As a result of the transplantation of a total of 302 injected embryos, 16 offspring mice were born. Chimerism in the offsprings can be determined by the extent of TT2 cell-derived agouti coat color (dark brown) in the host embryo-derived albino coat color. Out of the 16 produced offsprings, 5 mice were recognized to have a partial agouti coat color, indicating the contribution of the ES cell.

These results show that the ES cell clone PG22-5 retaining a human chromosome #22 partial fragment and a human chromosome #14 partial fragment maintains the ability to produce chimera, that is, the ability to differentiate into normal tissues of mouse.

25 Example 37

Detection of human antibody λ chain and μ chain in sera of the chimeric mice derived from the ES cells retaining both a human chromosome #22 partial fragment and a human chromosome #14 partial fragment

30 The chimeric mice KPG22-9, 10 and 12 from Example 36 were immunized with HSA. The chimeric mice KPG22-9 and 10 were immunized 11 weeks after birth and bled 2 weeks after the immunization. The chimeric mouse KPG22-12 was immunized twice at 7 and 11 weeks after birth and bled 2 weeks after the second immunization.

35 A serum human antibody μ chain, a serum human antibody λ chain, and a serum antibody having both human antibody λ and μ chains were detected by ELISA in accordance with Example 14.

For the detection of complete human antibody molecules, 96-well microtiter plates were coated with anti-human immunoglobulin λ chain antibody (Kirkegaard & Perry Laboratories Inc., 01-10-11) diluted with PBS and a serum sample was added. Subsequently, peroxidase-labeled anti-human immunoglobulin μ chain antibody (The Binding Site Limited, MP008) was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added as a 40 peroxidase substrate, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgM antibody having λ chain (Dainippon Pharmaceutical Co., Ltd, U13200) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS. Human antibody μ and λ chains were detected and determined quantitatively by ELISA in the same manner as in Examples 29 and 32. The results are shown in Table 14.

Table 14

Concentrations of Human Antibodies in Chimeric Mice (ELISA)					
ES clone	Chimeric mouse	Chimerism (%)	IgM (mg/l)	Ig λ (mg/l)	IgM, λ (mg/l)
PG22-5	KPG22-9	30	2.54	9.9	0.043
PG22-5	KPG22-10	5	4.96	21.5	0.333
PG22-5	KPG22-12	40	3.71	7.0	0.048
3-2	K9	50	6.66	-	< 0.003
PG22-1	KPG22-2	50	-	17.6	< 0.003

Both λ and μ chains were detected in the chimeric mice. An antibody molecule having both human antibody μ and λ chains was detected. These results show: the human antibody λ chain gene and human antibody μ chain gene can function at the same time in the chimeric mice derived from the ES cells retaining human chromosome #22 partial fragments and human chromosome #14 partial fragments; and a complete antibody containing both human heavy and light chains was produced in part of the B cells.

The control mice, that is, the chimeric mouse K9 retaining only human chromosome #14 from Example 10 and the chimeric mouse KG22-2 retaining only human chromosome #22 from Example 31, gave background levels of an antibody having both human antibody λ and μ chains in the sera. It was confirmed that in these detection systems, only a complete antibody molecule having human λ and μ chains was detected.

10

Example 38

Detection of human antibody having human κ and μ chains in sera of the chimeric mice derived from the ES cells retaining both human chromosome #2 partial fragments and human chromosome #14 partial fragments

15

The chimeric mouse KPG-15 (derived from the TT2ES clone PG5, 10% chimerism) was immunized during 2-3 months after birth 3 times with 0.2 ml of a solution of human serum albumin (HSA, Sigma, A3782) and adjuvant (MPL+TDM Emulsion, RIBI Immunochem Research Inc.) in PBS at a HSA concentration of 0.25 mg/ml and bled (see Example 15). The chimeric mouse KPG-26 (derived from the TT2ES clone PG6, 40% chimerism) of 6 weeks after birth was bled. The concentration of a complete human antibody molecule in the sera was determined by ELISA in accordance with Example 14. Ninety six-well microtiter plates were coated with anti-human immunoglobulin κ chain antibody (Kirkegaard & Perry Laboratories Inc., 01-10-10) diluted with PBS, and a serum sample was added. Subsequently, peroxidase-labeled anti-human immunoglobulin μ chain antibody (The Binding Site Limited, MP008) was added to the plates and incubated. After TMBZ (Sumitomo Bakelite, ML-1120T) was added as a peroxidase substrate, enzyme activity was determined by absorbance measurement at 450 nm. Purified human IgM antibody having κ chain (CAPPÉL, 6001-1590) was used as standard. The standard was diluted stepwise with mouse serum-supplemented PBS. The concentrations of κ chain and μ chain were determined by the same method as in Example 20. The results are shown in Table 15.

30

Table 15

Concentrations of Human Antibodies in Chimeric Mice (ELISA)					
ES clone	Chimeric mouse	Chimerism (%)	IgM (mg/l)	Ig κ (mg/l)	IgM, κ (mg/l)
PG-5	KPG15	10	0.18	1.01	0.075
PG-6	KPG26	40	1.52	1.26	0.018
3-2	K9	50	6.66	-	< 0.002
5-1	K2-9	40	-	135	< 0.002

A antibody molecule having both human antibody μ and κ chains was detected. The control mice, that is, the chimeric mouse K9 retaining only human chromosome #14 from Example 10 and the chimeric mouse K2-9 retaining only human chromosome #2 from Example 13, gave background levels (< 0.002 mg/ml) of an antibody having human antibody κ and μ chains in the sera. These results show: the human antibody κ chain gene and human antibody μ chain gene can function at the same time in the chimeric mice derived from the ES cells retaining both human chromosome #2 partial fragments and human chromosome #14 partial fragments; and a complete antibody molecule containing both human heavy and light chains was produced in part of the B cells.

50

Example 39

Preparation of mouse ES cell clone (TT2F, XO) retaining a human chromosome #2 partial fragment

The cell clone PG1 from Example 16 was used as a chromosome donor cell for the preparation of a mouse ES cell (XO) retaining a human chromosome #2 partial fragment. A TT2F cell (purchased from Lifetec Oriental Co.) having a karyotype of (39, XO), which was reported to differentiate efficiently into an oocyte in chimeric mice (Shinichi Aizawa, "Biomanual Series 8, Gene Targeting" published by Yodosha, 1995), was used as a chromosome recipient cell. The experiment of microcell fusion and the selection of puromycin resistant cells were carried out by the same methods as

in the selection of the G418 resistant clones in Example 9 except that the concentration of puromycin was 0.75 µg/ml. The frequency of the appearance of the puromycin resistant clones was 5 per 10^7 of TT2F cells. The puromycin resistant clones were stored frozen and genomic DNAs were prepared from the clones by the same methods as in Example 2. The retention of human chromosome #2 partial fragments in the drug resistant clones P-20 and P-21 was confirmed by PCR analysis. As a result of PCR amplification using genomic DNAs of the drug resistant clones as templates and three kinds of primers Ck, FABP1 and V_k 1-2 whose presence in the A9/#2 W23 clone was confirmed in Example 1, all of the three primers gave expected amplification products in both of the two clones.

These experiments demonstrate that the obtained puromycin resistant ES cell clone (TT2F, XO) retained a human chromosome #2 partial fragment.

10 Example 40

Production of the chimeric mice from the mouse ES cell clone (TT2F, XO) retaining a human chromosome #2 partial fragment

15 The cells in a frozen stock of the puromycin resistant TT2F cell clone P-21 from Example 39 which was confirmed to retain a human chromosome #2 partial fragment were thawed, started to culture and injected into 8-cell stage embryos obtained by mating ICR or MCH(ICR) male and female mice (CREA JAPAN, INC.); the injection rate was 10-12 cells per embryo. The embryos were cultured in a medium for ES cells (see Example 9) overnight to develop to blastocysts. Two and a half day after a foster mother ICR mouse was subjected to a pseudopregnant treatment, about ten 20 of the injected embryos were transplanted to each side of the uterus of the foster mother. The results are shown in Table 16.

25 Table 16

Production of the chimeric mice from the TT2F cell clone retaining a human chromosome #2 partial fragment								
ES cell clone/human chromosome	Puromycin resistant clone No.	Number of ES cell-injected 8-cell stage embryos	Number of offspring mice	Number of chimeric mice	Contribution to coat color			
					<20%	20-50%	50-90%	100%
TT2F/#2fg.	P-21	141	20	9	0	2	3	4

40 As a result of the transplantation of a total of 141 injected embryos, 20 offspring mice were born. Chimerism in the offsprings can be determined by the extent of TT2F cell-derived agouti coat color (dark brown) in the host embryo-derived albino coat color. Out of the 20 produced offsprings, 9 mice were recognized to have a partial agouti coat color, indicating the contribution of the ES cell. Four of the 9 mice were chimeric mice having a full agouti coat color from the ES cells.

45 These results show that the ES cell clone P-21 retaining a human chromosome #2 partial fragment maintains the ability to produce chimera, that is, the ability to differentiate into normal tissues of mouse.

Example 41

Detection and quantitative determination of human antibody κ chain in sera of the chimeric mice derived from the TT2F clone retaining a human chromosome #2 partial fragment

50 The chimeric mice K2-1F, 2F, 3F and 4F (derived from the P-21 clone, 100% chimerism) from Example 40 of about 1 month after birth were bled and the concentration of human antibody κ chain in the sera was determined quantitatively by ELISA in the same manner as in Example 20.

55 The results are shown in Table 17. It was confirmed that the human antibody κ chain gene could function in the chimeric mice when the TT2F was used as an ES cell.

Table 17

Concentration of Human Antibody κ Chain in Chimeric Mice (ELISA)		
Chimeric mouse	%Chimerism	Ig κ (mg/l)
K2-1F	100	66
K2-2F	100	156
K2-3F	100	99
K2-4F	100	20

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10

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Example 42

Confirmation of the retention of human chromosome in progenies of the chimeric mice derived from the mouse ES cell (TT2F, XO) retaining a human chromosome #2 partial fragment

Examination was made as to whether ES cell-derived progenies would be reproduced by mating the female chimeric mice K2-1F and K2-4F (both were of 100% chimerism in coat color) from Example 40 with ICR male mice. In such a mating, offspring mice of an agouti coat color should be reproduced from the oocytes derived from the TT2F cell (agouti coat color, dominant) in the chimeric mice and offspring mice of an albino coat color should be reproduced from oocytes derived from ICR if the oocytes are fertilized with the sperms from ICR male mice (albino, recessive). All the viable offspring mice (K2-1F, 10 mice and K2-4F, 5 mice) obtained by one mating of the respective combinations had an agouti coat color which derived from the ES cells. The retention of human chromosome fragments in genomic DNAs prepared from the tails of the offspring mice was examined by a PCR method. As a result of the PCR amplification using three kinds of primers whose presence in the P-21 clone (see Example 39) was confirmed, the presence of these three markers was confirmed in 4 out of the ten mice from K2-1F and in 2 out of the five mice from K2-4F. The results of the PCR of these 15 offspring mice are shown in Fig. 21. In Fig. 21, markers (ϕ X174/HaeIII fragment, Nippongene) and the DNA molecular weights of main bands are shown at the right side and the lengths of expected products of amplification with the respective primers are shown by arrows at the left side. At the right side, the results with tail-derived DNA of the mother chimeric mice K2-1F and K2-4F (positive controls) are shown. These results show that the TT2 cell clone P-21 differentiated into functional oocyte in the chimeric mice and that a human chromosome #2 partial fragment was transmitted to offsprings through oocyte.

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Example 43

Confirmation of the retention of human chromosome in progenies of the chimeric mice derived from the mouse ES cell (TT2, XY) retaining a human chromosome #2 partial fragment

Examination was made as to whether ES cell-contributed offspring mice would be produced by mating K2-18 (70% chimeric male mouse from Example 13) with K2-19 (60% chimeric female mouse of Example 13) or non-chimeric female littermates. Since TT2 cell has the karyotype of (40, XY), it may differentiate into a functional sperm in the male chimeric mouse K2-18. If this is the case, offspring mice of an agouti coat color should be reproduced from ICR (albino, recessive)-derived oocytes fertilized with sperms from TT2 cell (agouti color, dominant) in the chimeric mice. While a total of 110 viable offspring mice were obtained by the mating, ten had an agouti coat color which derived from the ES cells. The retention of human chromosome fragments in genomic DNAs prepared from the tails of 7 out of the ten offspring mice of an agouti coat color was examined by a PCR method. As a result of PCR amplification using two kinds of primers Cκ and FABP1 whose presence in the 5-1 clone (TT2/#2fg.. Example 12) was confirmed and primer Vκ 1-2 which was shown in Example 1, the presence of all of the three markers was confirmed in 2 out of the seven mice. These results show that the TT2 cell clone 5-1 retaining a human chromosome #2 partial fragment differentiated into functional sperms in the chimeric mice and that the human chromosome #2 partial fragment was transmitted to offsprings through the sperms.

Example 44

Detection and quantitative determination of human antibody κ chain in sera of offspring mice of the chimeric mice

5 The concentration of human antibody κ chain in the sera of the offspring mice K2-1F-1~10 and K2-4F-1~5 from Example 42 was determined quantitatively by ELISA. The mice of about 4~6 weeks after birth were bled and the concentration of human antibodyκ chain in the sera was determined by ELISA in the same manner as in Example 20.

10 The results are shown in Table 18 together with the data obtained in Example 42 on the retention of chromosome. It was confirmed that the human antibody κ chain gene can function in the offspring mice reproduced from the chimeric mice.

Table 18

Concentration of Human Antibody κ Chain in Offspring Mice (ELISA)				
	Mother mouse	Number of mouse	Presence of human chromosome #2 fragments	Ig κ (mg/l)
20	K2-1F	#1	-	0.58
	K2-1F	#2	+	84.1
	K2-1F	#3	+	12.8
	K2-1F	#4	+	15.1
	K2-1F	#5	-	0.52
	K2-1F	#6	-	0.58
	K2-1F	#7	-	1.30
	K2-1F	#8	-	0.90
	K2-1F	#9	-	0.56
	K2-1F	#10	+	28.8
30	K2-4F	#1	-	< 0.04
	K2-4F	#2	+	23.3
	K2-4F	#3	+	11.8
	K2-4F	#4	-	0.08
	K2-4F	#5	-	0.06

Example 45

45 Analysis of spleen cells from the human chromosome #14 partial fragment transferred chimeric mice

Flow cytometry analysis was accordance with the method described in "New Biochemical Experiment Lecture 12, molecular immunology I-Immunocells • Cytokines-", edited by the Japanese Biochemical Society, 1989, published by Tokyo Kagaku Dojin; "Cell Technology Separated Volume 8, New Cell Technology Experiment Protocol", edited by the University of Tokyo, Medical Science Institute, Anti-cancer Laboratory, 1991, published by Shujunsha; and A.Doyle and J.B.Griffiths, "Cell & Tissue Culture: Laboratory Procedures", published by John Wiley & Sons Ltd., 1996. The spleen was removed from the chimeric mouse KPG06 (derived from the PG16 clone, 30% chimerism) from Example 19 of six months after birth and treated with an aqueous solution of ammonium chloride. The spleen cells were stained with fluorescein isothiocyanate (FITC)-labeled anti-mouse CD45R (B220) antibody (Pharmingen, 01124A) in PBS containing 1% rat serum. After being washed, the cells were reacted with 0.1 µg of biotin-labeled anti-human IgM antibody (Pharmingen, 08072D) or a control biotin-labeled antihuman λ chain antibody (Pharmingen, 08152D) in PBS containing 5% mouse serum and stained with 0.1 µg of streptavidin-phycoerythrin (Pharmingen, 13025D), followed by analysis with a flowcytometer (Becton Dickinson Immunocytometry Systems, FACSsort). An ICR mouse retaining no human

chromosom was used as a control for analysis by the same method. The results are shown in Fig. 22. In Fig. 22, the human IgM is plotted on the horizontal axis and the CD45R (B220) is plotted on the vertical axis. A population of cells positive to both B cell marker CD45R (FITC) and human IgM (PE) increased by 4%, indicating that cells expressing human antibody μ chain on the cell surfaces were present in the chimeric mice.

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Example 46

Cloning and sequencing of variable regions of human antibody genes from cDNA derived from the spleen of the chimeric mice expressing human antibody heavy chain, κ and λ chains, respectively

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In the same manner as in Example 5, cDNAs were synthesized from RNAs extracted from the spleens of the chimeric mice K15A (derived from the 1-4 clone, prepared by the method described in Example 10), K2-8 prepared in Example 13 and KPG22-2 prepared in Example 31, all of which were confirmed to express human antibody heavy chain, κ and λ chains in Examples 29, 23 and 32, respectively. PCR was performed using the respective cDNAs and the following primers to amplify the variable regions of respective human antibody cDNA derived from the spleen of a non-chimeric mouse ICR was used as a negative control. Those primers set forth below without indication of reference literature were designed on the basis of the nucleotide sequences obtained from data bases such as Genebank and the like.

15

• K15A (heavy chain)

For constant region:

HGMEX1-2 : 5'-CCAAGCTTCAGGAGAAAGTGATGGAGTC (SEQ ID NO:53)

HGMEX1-1 : 5'-CCAAGCTTAGGCAGCCAACGGCCACGCT (used in 2nd PCR of VH3BACK) (SEQ ID NO:54)

20

For variable region:

VH1/5BACK (59°C, 35 cycles, Marks et al., Eur. J. Immunol., 21, 985-, 1991),

VH4BACK (59°C, 35 cycles, Marks et al., supra), and

VH3BACK (1st PCR:59 °C, 35 cycles; 2nd PCR:59°C, 35 cycles, Marks et al., supra)

25

• K2-8 (light chain κ)

For constant region:

KC2H: 5'-CCAAGCTTCAGAGGCAGTCCAGATTTC (SEQ ID NO:55)

For variable region:

30

Vk1/4BACK (55°C, 40 cycles, Marks et al., Eur. J. Immunol., 21, 985-, 1991),

Vk2BACK (55°C, 40 cycles, Marks et al., supra), and

Vk3BACK (55°C, 40 cycles, Marks et al., supra)

• KPG22-2 (light chain λ)

35

For constant region:

C λ MIX (a mixture of the following three kinds of primers at an equal molar ratio)

IGL1-CR : 5'-GGGAATTGGGTAGAACGTCACTGATCAG (SEQ ID NO:56)

IGL2-CR : 5'-GGGAATTGGGTAGAACGTCACTTATGAG (SEQ ID NO:57)

40

IGL7-CR : 5'-GGGAATTGGGTAGAACGTCACTTACGAG (SEQ ID NO:58)

For variable region:

V λ 1LEA1 (55°C, 40 cycles, Williams et al., Eur. J. Immunol., 23, 1456-, 1993),

V λ 2MIX (55°C, 40 cycles, a mixture of V λ 2 LEA1 and V λ 2 JLEAD (Williams et al. (supra)) at an equal molar ratio)

45

V λ 3MIX (55°C, 40 cycles, a mixture of V λ 3LEA1, V λ 3JLEAD and V λ 3BACK4, which were reported in Williams et al. (supra) at an equal molar ratio.

The PCR was performed with combinations of the primers for constant regions with those for variable regions (3 primer pairs each for heavy chain, κ and λ chains) at 94 °C for 15 seconds, at the annealing temperatures shown with respect to the respective primers for variable region for 15 seconds, at 72 °C for 20 seconds in the cycle numbers shown with respect to the respective primers for variable region using a Thermal Cycler 9600 (Perkin-Elmer Corp.). In the sec-

50

ond run of PCR using VH3BACK, the amplification products of the first run of PCR were amplified again with a combination of the two primers H1GMEX1-1 and VH3BACK. All of the amplification products were electrophoresed on a 1.5% agarose gel and stained with ethidium bromide for detection. As a result, the amplification products having expected lengths (heavy chain, about 470 bp; light chain κ , about 400 bp; and light chain λ , about 510 bp) were detected in all of the combinations. In the negative control, specific amplification product was not detected at the same position in any of

the combinations. The obtained amplification products were extracted from the agarose gel using prep.A.gene (Bio-Rad Laboratories, Inc.), treated with restriction enzymes (heavy chain, HindIII and PstI; light chain κ , HindIII and Pvull; and light chain λ , HindIII and EcoRI), and cloned into pUC119 (TAKARA SHUZO CO., LTD.) at the sites of HindIII/PstI (heavy chain), HindIII/HincII (κ chain) and HindIII/EcoRI (λ chain). The nucleotide sequences of the products that were amplified with the following primers and which were cloned into the plasmids were determined with a Fluorescence Autosequencer (Applied Biosystems Inc.).

- HIGMEX1-2×VH1/5BACK: 10 clones
- HIGMEX1-2×VH4BACK: 8 clones
- 10 • HIGMEX1-2 (2nd PCR, HIGMEX1-1)×VH3BACK: 5 clones
- KC2H×V κ 1/4BACK: 6 clones
- KC2H×V κ 2BACK: 7 clones
- KC2H×V κ 3BACK: 4 clones
- CAMIX×V λ .1LEA1: 5 clones
- 15 • CAMIX×V λ .2MIX: 6 clones
- CAMIX×V λ .3MIX: 5 clones

The obtained nucleotide sequences were analyzed with DNASIS (Hitachi Software Engineering Co., Ltd.). The results show that all of the sequences were derived from human and that they were functional sequences which did not contain a termination codon at any site between an initiation codon and a constant region: this was true with all of the κ and λ chains and with 21 out of a total of 23 heavy chains. When the same sequences were removed from the determined sequences, unique variable region sequences were identified. as follows: 17 heavy chains, 11 κ chains, and 12 λ chains.

25 Example 47

Analysis of the nucleotide sequences of variable region of human antibody genes from cDNA derived from the spleen of the chimeric mouse expressing human antibody heavy chain, κ and λ chains, respectively

30 The nucleotide sequences determined in Example 46 (heavy chain, 17 clones; κ chain, 11 clones; and λ chain, 12 clones) were analyzed in the following points.

1. Identification of known germ line V gene segments used in the respective variable regions
2. Identification of known germ line J gene segments used in the respective variable regions
- 35 3. Identification of known germ line D gene segments used in the heavy chain variable regions
4. Identification of the addition of N region in the heavy chain variable regions on the basis of the results of 1, 2 and 3
5. Determination of the amino acid sequences deduced from the nucleotide sequences of the respective variable regions

40 The results are shown in Table 19. For the identification in points of 1 and 2, search for homology with germ line V and J segments registered in Genbank and the like was conducted with DNASIS. The VH segments, V κ segments and V λ segments are shown in Table 19 together with the family names of the respective V fragments in accordance with the conventions described in Cook et al., Nature genetics, 7, 162-, 1994 (VH fragments), Klein et al., Eur. J. Immunol, 23, 3248-, 1993 (V κ fragments) and Williams et al. (supra) (V λ fragments), respectively. For the identification in point 3, search for homology with germ line D fragments reported in Ichihara et al., The EMBO J., 7, 13, 4141-, 1988 was conducted with DNASIS. Assignment was based on at least 8 bp identity and the results are shown in Table 19. DN1* is believed to be the new DN family segment reported in Green et al., Nature Genetics, 7, 13-, 1994. For the identification in point 4, the nucleotide sequences which did not appear in any germline sequences were determined to be N regions on the basis of the results for 1(V), 2(J) and 3 (D). As a result, N region was observed in 11 of the 13 sequences in which D segment was identified and its average length was 8.7 bp. For the determination in point 5, the respective sequences were converted by DNASIS to amino acid sequences which were expressed with one letter symbols. In Table 19, only CDR3 region is shown. At the right side of Table 19, the names of the primers used in cloning of the respective variable regions and the names of clones are shown.

Table 19

5

	V family	V segment	CDR3	J (D)	V primer	Clone
	K15A					
10	VH1	VH1-8	VRSSSWYEYYYYGMDV	J6 (DN1)	VH4BACK	H4-10
		VH1-18	GGITMVRGLIITDWYFDL	J2 (DXP'1)	VH1/5BACK	H1-7
		VH1-24	APYSGRFDY	J4 (DK1)	VH1/5BACK	H1-6
		VH1-46	ERYYCGSYQDYYYYGMDV	J6 (DXP'1)	VH1/5BACK	H1-2
		VH1-46	GGYSGYEDYYYYGMDV	J6 (DK1)	VH1/5BACK	H1-10
15	VH2	VH2-5	SYFDWPDFDY	J4 (DXP1)	VH4BACK	H4-14
	VH3	VH3-21	EGCSGGSCLPGYYYYGMDV	J6 (DLR2)	VH1/5BACK	H1-4
		VH3-23	AHGDPYFDY	J4	VH1/5BACK	H1-3
		VH3-23	DADAFDI	J3	VH1/5BACK	H1-8
		VH3-23	SGWDY	J4 (DN1*)	VH3BACK	H3-3
20		VH3-23	TGFDL	J2	VH4BACK	H4-4
		VH3-33	EGGYGSVGDDYYYYGMDV	J6 (DXP'1)	VH1/5BACK	H1-9
		VH3-33	GGYSYGYDYYYYGMDV	J6 (DXP'1)	VH3BACK	H3-5
		VH3-33	GYSSGWYDY	J4 (DN1*)	VH4BACK	H4-9
25	VH4	VH4-34	RYSSGWYYFDY	J4 (DN1*)	VH4BACK	H4-15
		VH4-59	GRIAVASF DY	J4 (DN1*)	VH4BACK	H4-2
		VH4-59	GSGSYFHF DY	J4	VH4BACK	H4-6
	K2-8					
	V κ 1	O18-8	QQHDNL PFT	J3	V κ 1BACK	K1-1
		O18-8	QQYDNL PIT	J5	V κ 1BACK	K1-3
30		O18-8	QQHDNL PFA	J3	V κ 2BACK	K2-2
		L1	QQYN SYPLT	J4	V κ 1BACK	K1-6
	V κ 2	A17	MQGTHLLT	J4	V κ 2BACK	K2-1
		A17	MQGTHWIT	J5	V κ 2BACK	K2-5
35	V κ 3	A27	QQYGSSPTWT	J1	V κ 3BACK	K3-1
		A27	QQYGSSPFT	J3	V κ 3BACK	K3-4
		A27	QQYGSSPLWT	J1	V κ 3BACK	K3-5
		A27	QQYGSSPPWT	J1	V κ 3BACK	K3-6
	V κ 6	A26-10	HQS SLPQT	J1	V κ 2BACK	K2-4
	KPG22-2					
40	V λ 1	DPL3	AAWDDSLDV V	JC3	V λ 1LEA1	L1-3
		DPL5	GTWDSSL SAGV	JC2	V λ 1LEA1	L1-4
		DPL5	GTWDSSL SAGVV	JC3	V λ 1LEA1	L1-6
		DPL5	GTWDSSL SAVV	JC2	V λ 1LEA1	L1-9
45	V λ 2	DPL8	QSYDSSL SGVV	JC3	V λ 1LEA1	L1-8
		DPL10	CSYAGSSTLV	JC2	V λ 2MIX	L2-4
		DPL11	SSYTSSSTV V	JC2	V λ 2MIX	L2-1
		DPL11	SSYTSSSTLV	JC2	V λ 2MIX	L2-3
		DPL11	CSYTSSSTFV	JC2	V λ 2MIX	L2-7
50		DPL12	SSYAGSNNLV	JC3	V λ 2MIX	L2-5
		DPL12	SSYAGSNNFV V	JC3	V λ 2MIX	L2-6
	V λ 3	DPL16	NSRDSSGNLV	JC2	V λ 3MIX	L3-1

55

INDUSTRIAL EXPLOITABILITY

In accordance with the present invention, chimeric non-human animals that retain foreign chromosomes or frag-

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ments thereof and which express the genes on the foreign chromosomes or fragments, as well as their progenies can be produced. The chimeric non-human animals and their progenies can be used to produce biologically active substances.

5

SEQUENCE LISTING

SEQ ID NO: 1

10

SEQUENCE LENGTH: 20

15

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

20

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

25

TGGAAGGTGG ATAACGCCCT

20

SEQ ID NO: 2

25

SEQUENCE LENGTH: 22

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

30

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

35

TCATTCTCCT CCAACATTAG CA

22

SEQ ID NO: 3

40

SEQUENCE LENGTH: 21

SEQUENCE TYPE: nucleic acid

45

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

50

SEQUENCE DESCRIPTION

TGTAGGGGAC CTGGAGCCTT G

21

55

SEQ ID NO: 4

SEQUENCE LENGTH: 21
SEQUENCE TYPE: nucleic acid
5 STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
10 SEQUENCE DESCRIPTION
TTGACAACTC ACCTGGACTA G
15 SEQ ID NO: 5
SEQUENCE LENGTH: 20
20 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
25 MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
CTCTCCTGCA GGGCCAGTCA
30 SEQ ID NO: 6
SEQUENCE LENGTH: 22
35 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
40 MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
TGCTGATGGT GAGAGTGAAC TC
45 SEQ ID NO: 7
SEQUENCE LENGTH: 20
50 SEQUENCE TYPE: nucleic acid

55

STRANDEDNESS: single
5 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
10 AGTCAGGGCA TTAGCAGTGC 20

SEQ ID NO: 8
15 SEQUENCE LENGTH: 20
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
20 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
25 GCTGCTGATG GTGAGAGTGA 20

SEQ ID NO: 9
30 SEQUENCE LENGTH: 20
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
35 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
40 TGGTGGCTGA AAGCTAAGAA 20

SEQ ID NO: 10
45 SEQUENCE LENGTH: 20
SEQUENCE TYPE: nucleic acid
50 STRANDEDNESS: single
TOPOLOGY: linear

55

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

5 CCAGAAAGAAT GGTGTCATTA

20

10 SEQ ID NO: 1 1

SEQUENCE LENGTH: 2 0

SEQUENCE TYPE: nucleic acid

15 STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

20 SEQUENCE DESCRIPTION

TCCAGGTTCT GCAGAGCAAG

20

25 SEQ ID NO: 1 2

SEQUENCE LENGTH: 2 0

SEQUENCE TYPE: nucleic acid

30 STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

35 SEQUENCE DESCRIPTION

TGTAGTTGGA GGCCATGTCC

20

40

SEQ ID NO: 1 3

SEQUENCE LENGTH: 2 0

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

50 MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

55

CCCCACCCAT GATCCAGTAC

20

5

SEQ ID NO: 1 4

SEQUENCE LENGTH: 2 0

10

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

15

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

GCCCTCAGAA GACGAAGCAG

20

20

SEQ ID NO: 1 5

SEQUENCE LENGTH: 2 2

25

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

30

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

35

GAGAGTTGCA GAAGGGGTGA CT

22

40

SEQ ID NO: 1 6

SEQUENCE LENGTH: 2 2

SEQUENCE TYPE: nucleic acid

45

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

50

SEQUENCE DESCRIPTION

GGAGACCACC AAACCCCTCCA AA

22

55

SEQ ID NO: 1 7
SEQUENCE LENGTH: 2 0
5 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
10 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
15 GGCTATGGGG ACCTGGGCTG 20

SEQ ID NO: 1 8
SEQUENCE LENGTH: 2 2
20 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
25 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
30 CAGAGACACA GGCACGTAGA AG 22

SEQ ID NO: 1 9
35 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
40 STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
45 SEQUENCE DESCRIPTION
TTAAGGGTCA CCCAGAGACT 20

50 SEQ ID NO: 2 0
SEQUENCE LENGTH: 2 0

SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
5 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
10 SEQUENCE DESCRIPTION
TGTAGTTGGA GGCCATGTCC 20

15 SEQ ID NO: 2 1
SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
20 STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
25 SEQUENCE DESCRIPTION
CAAAAAGTCC AACCTATCA 20

30 SEQ ID NO: 2 2
SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
35 STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
40 SEQUENCE DESCRIPTION
GCCCTCAGAA GACGAAGCAG 20

45 SEQ ID NO: 2 3
SEQUENCE LENGTH: 2 0
50 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single

55

TOPOLOGY: linear
5 MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
TCGTTCCCTGT CGAGGATGAA 20
10 SEQ ID NO: 2 4
SEQUENCE LENGTH: 2 0
15 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
20 MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
TCACTCCGAA GCTGCCTTTC 20
25 SEQ ID NO: 2 5
SEQUENCE LENGTH: 2 1
30 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
35 MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
40 ATGTACAGGA TGCAACTCCT G 21
SEQ ID NO: 2 6
45 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
50 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
55

SEQUENCE DESCRIPTION
5 TCATCTGTAA ATCCAGGAGT 20

SEQ ID NO: 2 7
10 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
15 STRANDEDNESS: single
TOPOLOGY: linear
20 MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION
25 GATCCCATCG CAGCTACCGC 20

SEQ ID NO: 2 8
25 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
30 STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
35 SEQUENCE DESCRIPTION
35 TTTCGCCGAGT AGTCGCACGG 20

SEQ ID NO: 2 9
40 SEQUENCE LENGTH: 2 2
SEQUENCE TYPE: nucleic acid
45 STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
50 SEQUENCE DESCRIPTION
50 GATGAACTAG TCCAGGTGAG TT 22

SEQ ID NO: 3 0
5 SEQUENCE LENGTH: 2 2
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
10 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
15 CCTTTGGCT TCTACTCCTT CA 22

SEQ ID NO: 3 1
20 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
25 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
30 ATAGAGGGTA CCCACTCTGG 20

SEQ ID NO: 3 2
35 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
40 STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
45 AACCAAGGTAG GTTGATATGG 20

SEQ ID NO: 3 3
50

SEQUENCE LENGTH: 2 0
5 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
10 MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
AAGTTCCCTGT GATGTCAAGC 20
15 SEQ ID NO: 3 4
SEQUENCE LENGTH: 2 0
20 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
25 MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
TCATGAGGAG ATTAAACCCG 20
30 SEQ ID NO: 3 5
SEQUENCE LENGTH: 2 0
35 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
40 MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
TGTGAAGGAG GACCAGGTGT 20
45 SEQ ID NO: 3 6
50 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
55

STRANDEDNESS: single
5 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
10 TGTAGGGGTT GACAGTGACA 20

SEQ ID NO: 3 7
15 SEQUENCE LENGTH: 20
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
20 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
25 CTGAGAGATG CCTCTGGTGC 20

SEQ ID NO: 3 8
30 SEQUENCE LENGTH: 20
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
35 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
40 SEQUENCE DESCRIPTION
GGCGGTTAGT GGGGTCTTCA 20

45 SEQ ID NO: 3 9
50 SEQUENCE LENGTH: 20
SEQUENCE TYPE: nucleic acid
50 STRANDEDNESS: single
TOPOLOGY: linear
55

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

GGTGTCGTGG AACTCAGGCCG

20

5

SEQ ID NO: 4 0

SEQUENCE LENGTH: 20

SEQUENCE TYPE: nucleic acid

15

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

20

SEQUENCE DESCRIPTION

CTGGTGCAGG ACGGTGAGGA

20

25

SEQ ID NO: 4 1

SEQUENCE LENGTH: 20

SEQUENCE TYPE: nucleic acid

30

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

35

SEQUENCE DESCRIPTION

GCATCCTGAC CGTGTCCGAA

20

40

SEQ ID NO: 4 2

SEQUENCE LENGTH: 20

SEQUENCE TYPE: nucleic acid

45

STRANDEDNESS: single

TOPOLOGY: linear

50

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

55

GGGTCAGTAG CAGGTGCCAG

20

5

SEQ ID NO: 4 3

SEQUENCE LENGTH: 2 0

10

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

15

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

AGTGAGATAA GCAGTGGATG

20

20

SEQ ID NO: 4 4

SEQUENCE LENGTH: 2 0

25

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

30

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

35

GTTGTGCTAC TCCCATCACT

20

SEQ ID NO: 4 5

40

SEQUENCE LENGTH: 2 1

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

45

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

50

SEQUENCE DESCRIPTION

TTGTATTTCC AGGAGAAAGT G

20

55

SEQ ID NO: 4 6
SEQUENCE LENGTH: 2 0
5 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
10 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
15 GGAGACGAGG GGGAAAAGGG 20

SEQ ID NO: 4 7
SEQUENCE LENGTH: 2 7
20 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
25 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
30 ATGGACTGGA CCTGGAGGRT CYTCTKC 27

SEQ ID NO: 4 8
SEQUENCE LENGTH: 2 7
35 SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
40 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
45 SEQUENCE DESCRIPTION
ATGGAGYTTG GGCTGASCTG GSTTTYT 27

50 SEQ ID NO: 4 9
SEQUENCE LENGTH: 2 7
55

SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
5 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
10 SEQUENCE DESCRIPTION
ATGRAMMWAC TKTGKWCWY SCTYCTG 27
SEQ ID NO: 5 0
15 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
20 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
SEQUENCE DESCRIPTION
25 CAGAGGCAGT TCCAGATTTC 20
SEQ ID NO: 5 1
30 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
STRANDEDNESS: single
35 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid synthetic DNA
40 SEQUENCE DESCRIPTION
TGGGATAGAA GTTATTCA 20
SEQ ID NO: 5 2
45 SEQUENCE LENGTH: 2 0
SEQUENCE TYPE: nucleic acid
50 STRANDEDNESS: single
TOPOLOGY: linear
55

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

5 ATGGACATGR RRDYCCHVGY KCASCTT

27

10 SEQ ID NO: 5 3

SEQUENCE LENGTH: 2 8

SEQUENCE TYPE: nucleic acid

15 STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

20 SEQUENCE DESCRIPTION

CCAAGCTTCA GGAGAAAGTG ATGGAGTC

25 SEQ ID NO: 5 4

SEQUENCE LENGTH: 2 8

SEQUENCE TYPE: nucleic acid

30 STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

35 SEQUENCE DESCRIPTION

CCAAAGCTTAG GCAGCCAACG GCCACGCT

40

SEQ ID NO: 5 5

SEQUENCE LENGTH: 2 8

45 SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

50 MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

55

CCAAGCTTCA GAGGCAGTTC CAGATTTC

5

SEQ ID NO: 5 6

SEQUENCE LENGTH: 2 8

10

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

15

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

20

GGGAATTCGG GTAGAAGTCA CTGATCAG

SEQ ID NO: 5 7

25

SEQUENCE LENGTH: 2 8

SEQUENCE TYPE: nucleic acid

STRANDEDNESS: single

30

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

SEQUENCE DESCRIPTION

35

GGGAATTCGG GTAGAAGTCA CTTATGAG

40

SEQ ID NO: 5 8

SEQUENCE LENGTH: 2 8

SEQUENCE TYPE: nucleic acid

45

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid synthetic DNA

50

SEQUENCE DESCRIPTION

GGGAATTCGG GTAGAAGTCA CTTACGAG

55

SEQUENCE LISTING

5

(1) GENERAL INFORMATION:

(i) APPLICANT:

10

- (A) NAME: Kirin Beer Kabushiki Kaisha
- (B) STREET: 10-1, Shinkawa 2-chome
- (C) CITY: Chuo-ku
- 15 (D) STATE: Tokyo 104
- (E) COUNTRY: Japan
- (F) POSTAL CODE (ZIP): None

15

20 (ii) TITLE OF INVENTION: Chimeric animal and method for producing the
same

25

(iii) NUMBER OF SEQUENCES: 58

(iv) COMPUTER READABLE FORM:

30

- (A) MEDIUM TYPE: Floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)

35

(v) CURRENT APPLICATION DATA:

APPLICATION NUMBER: EP 96 113 839.3

40

(2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

45

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

50

(ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

5

TGGAAGGTGG ATAACGCCCT

20

(2) INFORMATION FOR SEQ ID NO: 2:

10

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

20

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

30

TCATTCTCCT CCAACATTAG CA

22

(2) INFORMATION FOR SEQ ID NO: 3:

35

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

40

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

50

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

5 GCAATCGGTC TGCCGGAAAGA

20

(2) INFORMATION FOR SEQ ID NO: 4:

10 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- 15 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

30 TTGGATCACT TTGGACCCAG

20

(2) INFORMATION FOR SEQ ID NO: 5:

35 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- 40 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

45 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic peptide"

50

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

5 CTCTCCTGCA GGGCCAGTCA

20

(2) INFORMATION FOR SEQ ID NO: 6:

10 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 22 base pairs
(B) TYPE: nucleic acid
15 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid
25 (A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

30 TGCTGATGGT GAGAGTGAAC TC

22

(2) INFORMATION FOR SEQ ID NO: 7:

35 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
40 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

45 (ii) MOLECULE TYPE: other nucleic acid
46 (A) DESCRIPTION: /desc = "synthetic DNA"

50

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

5 AGTCAGGGCA TTAGCAGTGC

20

(2) INFORMATION FOR SEQ ID NO: 8:

10 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
15 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

25 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

30 GCTGCTGATG GTGAGAGTGA

20

(2) INFORMATION FOR SEQ ID NO: 9:

35 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
40 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid
45 (A) DESCRIPTION: /desc = "synthetic DNA"

50

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

5 TGGTGGCTGA AAGCTAAGAA

20

(2) INFORMATION FOR SEQ ID NO: 10:

10 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- 15 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

30 CCAGAAGAAC GGTGTCATTA

20

(2) INFORMATION FOR SEQ ID NO: 11:

35 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- 40 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

45 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

50

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

55

TCCAGGTTCT GCAGAGCAAG

20

5 (2) INFORMATION FOR SEQ ID NO: 12:

(i) SEQUENCE CHARACTERISTICS:

- 10 (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

15 (ii) MOLECULE TYPE: other nucleic acid
20 (A) DESCRIPTION: /desc = "synthetic DNA"

25 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

TGTAGTTGGA GGCCATGTCC

20

30 (2) INFORMATION FOR SEQ ID NO: 13:

(i) SEQUENCE CHARACTERISTICS:

- 35 (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

40 (ii) MOLECULE TYPE: other nucleic acid
45 (A) DESCRIPTION: /desc = "synthetic DNA"

50 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:

55

CCCCACCCAT GATCCAGTAC

20

5

(2) INFORMATION FOR SEQ ID NO: 14:

(i) SEQUENCE CHARACTERISTICS:

10

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

15

(ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

20

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

GCCCTCAGAA GACGAAGCAG

20

30

(2) INFORMATION FOR SEQ ID NO: 15:

(i) SEQUENCE CHARACTERISTICS:

35

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

40

(ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

45

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:

GAGAGTTGCA GAAGGGGTGA CT

22

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(2) INFORMATION FOR SEQ ID NO: 16:

5 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

10

(ii) MOLECULE TYPE: other nucleic acid

15 (A) DESCRIPTION: /desc = "synthetic DNA"

20

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

25 GGAGACCACC AAACCCCTCCA AA

22

(2) INFORMATION FOR SEQ ID NO: 17:

30 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

35

(ii) MOLECULE TYPE: other nucleic acid

40 (A) DESCRIPTION: /desc = "synthetic DNA"

45

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

50 GGCTATGGGG ACCTGGGCTG

20

55

(2) INFORMATION FOR SEQ ID NO: 18:

5

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- 10 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

10

(ii) MOLECULE TYPE: other nucleic acid

15

(A) DESCRIPTION: /desc = "synthetic DNA"

20

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:

25

CAGAGACACA GGCACGTAGA AG

22

(2) INFORMATION FOR SEQ ID NO: 19:

30

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- 35 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

35

(ii) MOLECULE TYPE: other nucleic acid

40

(A) DESCRIPTION: /desc = "synthetic DNA"

45

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:

50

TTAAGGGTCA CCCAGAGACT

20

(2) INFORMATION FOR SEQ ID NO: 20:

55

5 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

10 (ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:

20 TGTAGTTGGA GGCCATGTCC

20

25 (2) INFORMATION FOR SEQ ID NO: 21:

30 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

35 (ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

40 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:

45 CAAAAAGTCC AACCTATCA

20

50 (2) INFORMATION FOR SEQ ID NO: 22:

55

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

5

10 (ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

15

20 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:

20

GCCCTCAGAA GACGAAGCAG

20

25

(2) INFORMATION FOR SEQ ID NO: 23:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

35

(ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

40

45 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

45

TCGTTCCCTGT CGAGGATGAA

20

50

(2) INFORMATION FOR SEQ ID NO: 24:

(i) SEQUENCE CHARACTERISTICS:

55

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

5

(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

10

15

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:

20

TCACTCCGAA GCTGCCTTTC

20

(2) INFORMATION FOR SEO ID NO: 25:

25

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 base pair
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

35

40

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:

45

ATGTACAGGA TGCAACTGCT G

21

(2) INFORMATION FOR SEQ ID NO: 26:

50

(i) SEQUENCE CHARACTERISTICS:

55

5
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

10
(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

15
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 26:

20
TCATCTGTAA ATCCAGCAGT

20

25
(2) INFORMATION FOR SEQ ID NO: 27:

25
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
30
(D) TOPOLOGY: linear

35
(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

40
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 27:

45
GATCCCACATCG CAGCTACCCG

20

(2) INFORMATION FOR SEQ ID NO: 28:

50
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid

55

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

5

(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

10

15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 28:

TTCGCCGAGT AGTCGCACGG

20

20 (2) INFORMATION FOR SEQ ID NO: 29:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 22 base pairs
25 (B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

30

(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

35

40 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 29:

GATGAACTAG TCCAGGTGAG TT

22

45 (2) INFORMATION FOR SEQ ID NO: 30:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 22 base pairs
50 (B) TYPE: nucleic acid
(C) STRANDEDNESS: single

55

(D) TOPOLOGY: linear

5 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

10

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 30:

15

CCTTTTGGCT TCTACTCCTT CA

22

20 (2) INFORMATION FOR SEQ ID NO: 31:

25

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

30

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

35

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 31:

40

ATAGAGGGTA CCCACTCTGG

20

45

(2) INFORMATION FOR SEQ ID NO: 32:

50

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

55

(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

5

10

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 32:

AACCAGGTAG GTTGATATGG

20

15

(2) INFORMATION FOR SEQ ID NO: 33:

20

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

25

(iii) MOLECULE TYPE: other nucleic acid

30

(A) DESCRIPTION: /desc = "synthetic DNA"

35

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 33:

40

AAGTTCCCTGT GATGTCAAGC

20

45

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

5

10 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 34:

TCATGAGCAG ATTAAACCCG

20

15

(2) INFORMATION FOR SEQ ID NO: 35:

20 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

25

(ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

30

35 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 35:

TGTGAAGGAG GACCAGGTGT

20

40

(2) INFORMATION FOR SEQ ID NO: 36:

45 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

50

(ii) MOLECULE TYPE: other nucleic acid

55

(A) DESCRIPTION: /desc = "synthetic DNA"

5

10 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 36:

TGTAGGGGTT GACAGTGACA

20

15 (2) INFORMATION FOR SEQ ID NO: 37:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

25

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

30

35 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 37:

CTGAGAGATG CCTCTGGTGC

20

40 (2) INFORMATION FOR SEQ ID NO: 38:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

50

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 38:

5 GGCGGTTAGT GGGGTCTTCA

20

(2) INFORMATION FOR SEQ ID NO: 39:

10 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
15 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 39:

30 GGTGTCGTGG AACTCAGGCG

20

(2) INFORMATION FOR SEQ ID NO: 40:

35 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
40 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid
45 (A) DESCRIPTION: /desc = "synthetic DNA"

50

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 40:

5 CTGGTGCAGG ACGGTGAGGA

20

(2) INFORMATION FOR SEQ ID NO: 41:

10 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
15 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 41:

30 GCATCCTGAC CGTGTCCGAA

20

(2) INFORMATION FOR SEQ ID NO: 42:

35 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
40 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

45 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

50

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 42:

5 GGGTCAGTAG CAGGTGCCAG

20

(2) INFORMATION FOR SEQ ID NO: 43:

10 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

15 20 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 43:

30 AGTGAGATAA GCAGTGGATG

20

(2) INFORMATION FOR SEQ ID NO: 44:

35 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

40 45 (ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "synthetic DNA"

50

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 44:

5 GTTGTGCTAC TCCCCATCACT

20

(2) INFORMATION FOR SEQ ID NO: 45:

10 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 21 base pairs
(B) TYPE: nucleic acid
15 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 45:

30 TTGTATTTCC AGGAGAAAGT G

21

(2) INFORMATION FOR SEQ ID NO: 46:

35 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
40 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

45 (ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

50

55

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 46:

5 GGAGACGAGG GGGAAAAGGG

20

(2) INFORMATION FOR SEQ ID NO: 47:

10 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 27 base pairs
(B) TYPE: nucleic acid
15 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: other nucleic acid
20 (A) DESCRIPTION: /desc = "synthetic DNA"

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 47:

30 ATGGACTGGA CCTGGAGGRT CYTCTKC

27

(2) INFORMATION FOR SEQ ID NO: 48:

35 (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 27 base pairs
(B) TYPE: nucleic acid
40 (C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid
45 (A) DESCRIPTION: /desc = "synthetic DNA"

50

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 48:

55

ATGGAGYTTG GGCTGASCTG GSTTTYT

27

5 (2) INFORMATION FOR SEQ ID NO: 49:

(i) SEQUENCE CHARACTERISTICS:

10 (A) LENGTH: 27 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

15 (ii) MOLECULE TYPE: other nucleic acid
20 (A) DESCRIPTION: /desc = "synthetic peptide"

25 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 49:

ATGRAMMWAC TKTGKWCWY SCTYCTG

27

30 (2) INFORMATION FOR SEQ ID NO: 50:

(i) SEQUENCE CHARACTERISTICS:

35 (A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

40 (ii) MOLECULE TYPE: other nucleic acid
45 (A) DESCRIPTION: /desc = "synthetic DNA"

50 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 50:

55

CAGAGGCAGT TCCAGATTTC

20

5 (2) INFORMATION FOR SEQ ID NO: 51:

(i) SEQUENCE CHARACTERISTICS:

10 (A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

15 (ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

20

25 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 51:

TGGGATAGAA GTTATTCAGC

20

30 (2) INFORMATION FOR SEQ ID NO: 52:

(i) SEQUENCE CHARACTERISTICS:

35 (A) LENGTH: 27 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

40 (ii) MOLECULE TYPE: other nucleic acid
(A) DESCRIPTION: /desc = "synthetic DNA"

45

50 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 52:

ATGGACATGR RRDYCCHVGY KCASCTT

27

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EP 0 773 288 A2

(2) INFORMATION FOR SEQ ID NO: 53:

5 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- 10 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

15 (A) DESCRIPTION: /desc = "synthetic DNA"

20

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 53:

25 CCAAGCTTCA GGAGAAAAGTG ATGGAGTC

28

(2) INFORMATION FOR SEQ ID NO: 54:

30 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- 35 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

40 (A) DESCRIPTION: /desc = "synthetic DNA"

45

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 54:

50 CCAAGCTTAG GCAGCCAACG GCCACGCT

28

55

EP 0 773 288 A2

(2) INFORMATION FOR SEQ ID NO: 55:

5 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

10 (ii) MOLECULE TYPE: other nucleic acid

15 (A) DESCRIPTION: /desc = "synthetic DNA"

20

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 55:

25 CCAAGCTTCA GAGGCAGTTC CAGATTTC

28

(2) INFORMATION FOR SEQ ID NO: 56:

30 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

35 (ii) MOLECULE TYPE: other nucleic acid

40 (A) DESCRIPTION: /desc = "synthetic DNA"

45

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 56:

50 GGGAAATTCGG GTAGAAGTCA CTGATCAG

28

(2) INFORMATION FOR SEQ ID NO: 57:

55

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

5

10 (ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

15

20 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 57:

20

GGGAATTCGG GTAGAAGTCA CTTATGAG

28

25

(2) INFORMATION FOR SEQ ID NO: 58:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 28 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

30

35 (ii) MOLECULE TYPE: other nucleic acid

- (A) DESCRIPTION: /desc = "synthetic DNA"

40

45 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 58:

45

GGGAATTCGG GTAGAAGTCA CTTACGAG

28

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Claims

1. A method for producing a chimeric non-human animal, which comprises preparing a microcell containing a foreign chromosome(s) or a fragment(s) thereof and transferring the foreign chromosome(s) or fragment(s) into a pluripotent cell by fusion with said microcell.
2. A method for producing a pluripotent cell containing a foreign chromosome(s) or a fragment(s) thereof, which comprises preparing a microcell containing a foreign chromosome(s) or a fragment(s) thereof and transferring the for-

sign chromosome(s) or fragment(s) into a pluripotent cell by fusion with said microcell.

3. The method of claim 1 or 2, wherein the foreign chromosome(s) or a fragment(s) thereof have a length(s) of at least 1 Mb.
5
4. The method of claim 1 or 2, wherein the foreign chromosome or fragment thereof contain a region encoding an antibody.
5. The method of claim 1 or 2, wherein the microcell containing a foreign chromosome(s) or a fragment(s) thereof is induced from a hybrid cell prepared by the fusion of a cell from which the foreign chromosome(s) or fragment(s) thereof is derived, with a cell having a high ability to form a microcell.
10
6. The method of claim 5, wherein the microcell containing a foreign chromosome(s) or a fragment(s) thereof is induced from a cell prepared by a further fusion of the microcell induced from said hybrid cell with a cell having a high ability to form a microcell.
15
7. The method of claim 5, wherein the cell from which the foreign chromosome(s) or fragment(s) thereof is derived is a human normal diploid cell.
- 20 8. The method of any one of claims 5-7, wherein the cell having a high ability to form a microcell is a mouse A9 cell.
9. The method of claim 1 or 2, wherein the pluripotent cell is an ES cell.
10. The method of claim 1, wherein the chimeric non-human animal is a mammal.
25
11. The method of claim 10, wherein the mammal is a mouse.
12. A pluripotent cell which is obtainable by the method of claim 2, said cell containing a foreign chromosome(s) or a fragment(s) thereof.
30
13. Use of the pluripotent cell of claim 12 for producing a chimeric non-human animal.
14. A chimeric non-human animal which is obtainable by the method of any one of claims 1 and 3-11, said animal retaining a foreign chromosome(s) or a fragment(s) thereof and expressing the gene(s) on the foreign chromosome(s) or fragment(s), or its progeny.
35
15. A tissue from the chimeric non-human animal or its progeny of claim 14.
16. A cell from the chimeric non-human animal or its progeny of claim 14.
40
17. The cell of claim 16, which is a B cell.
18. A hybridoma prepared by the fusion of the cell of claim 17 with a myeloma cell.
- 45 19. A method for producing a biologically active substance, which comprises expressing the gene on a foreign chromosome or a fragment in the chimeric non-human animal or its progeny of claim 14, or a tissue or a cell thereof, and recovering the biologically active substance as the expression product.
20. The method of claim 19, wherein the cell of the non-human animal or its progeny is a B cell.
50
21. The method of claim 20, wherein the B cell is immortalized by fusion with a myeloma cell.
22. The method of any one of claims 19-21, wherein the biologically active substance is an antibody.
- 55 23. The method of claim 22, wherein the antibody is an antibody of a mammal.
24. The method of claim 23, wherein the mammal is a human.

MICROCELL HYBRID CLONES											
	B-5	B-24	C group	D-1	D-2	D-8	D-14	E-21	I-23	L-34	M-16
S207	X	X	X	X	X	X	X	X	X	X	X
S177	X	X	X	X	X	X	X	X	X	X	X
POLYMORPHIC AND GENETIC MARKERS	●	●	●	●	●	●	●	●	●	●	●
Vk1-2 V _k 3-2	●	●	●	●	●	●	●	●	●	●	●
Ck	●	●	●	●	●	●	●	●	●	●	●
S156	X	X	X	X	X	X	X	X	X	X	X
S159	X	X	X	X	X	X	X	X	X	X	X
CHROMOSOME	I	I	T	I	I	I	I	I	I	T	I

N2 and
others
from
group N

T-9

W-23

Y-68

Z-15

I: INDEPENDENT T: TRANSLOCATED

FIG. 1 PCR ANALYSIS OF A9 CELLS RETAINING HUMAN CHROMOSOME #2 (FRAGMENT)

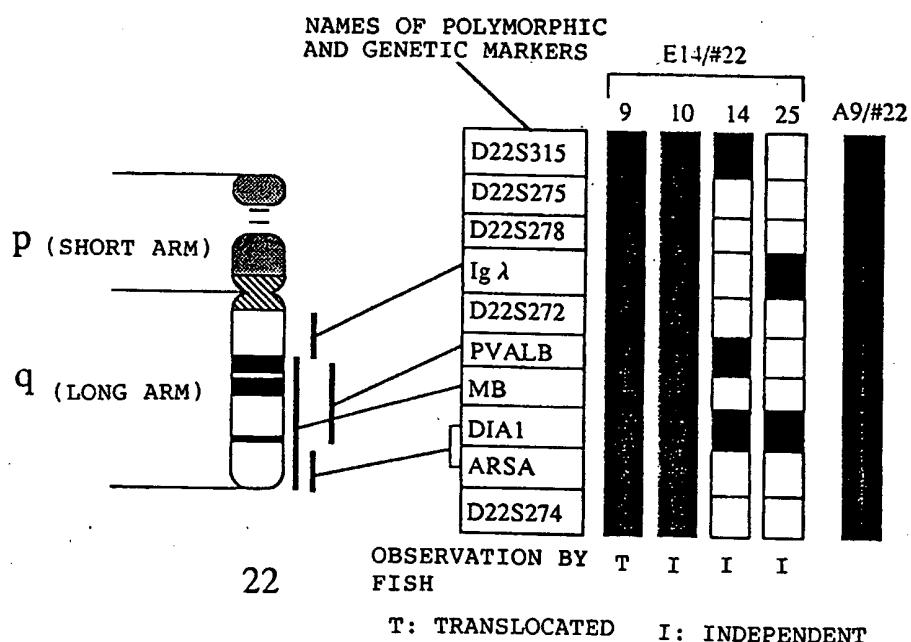


FIG. 2 RETENTION OF HUMAN CHROMOSOME #22 (FRAGMENTS)
IN E14 DRUG RESISTANT CLONES (PCR ANALYSIS)

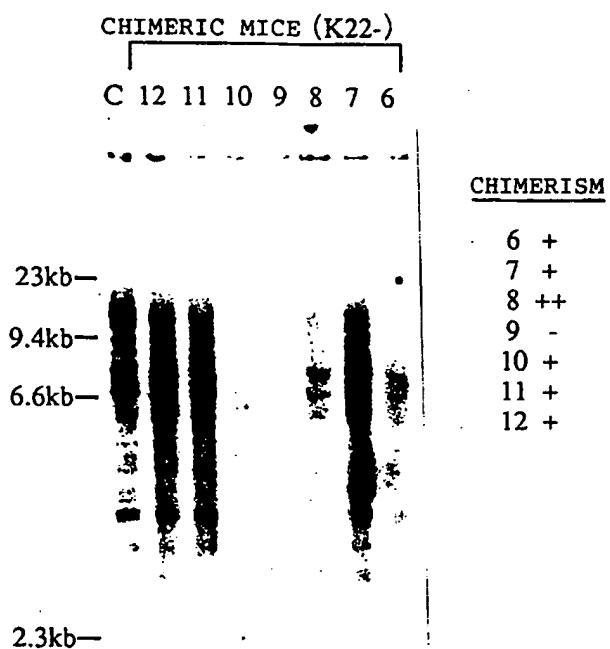


FIG. 3

RETENTION OF HUMAN LI SEQUENCE IN CHIMERIC MICE DERIVED
FROM CHROMOSOME #22 TRANSFERRED ES CELLS (SOUTHERN ANALYSIS)

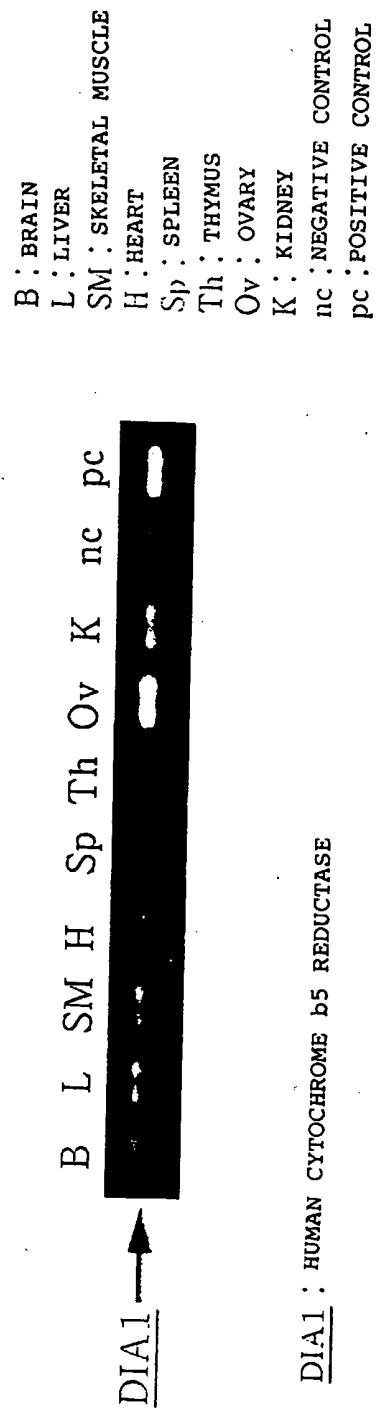


FIG. 4 RETENTION OF HUMAN CHROMOSOME IN ORGANS OF HUMAN CHROMOSOME #22-TRANSFERRED CHIMERIC MOUSE (PCR ANALYSIS)

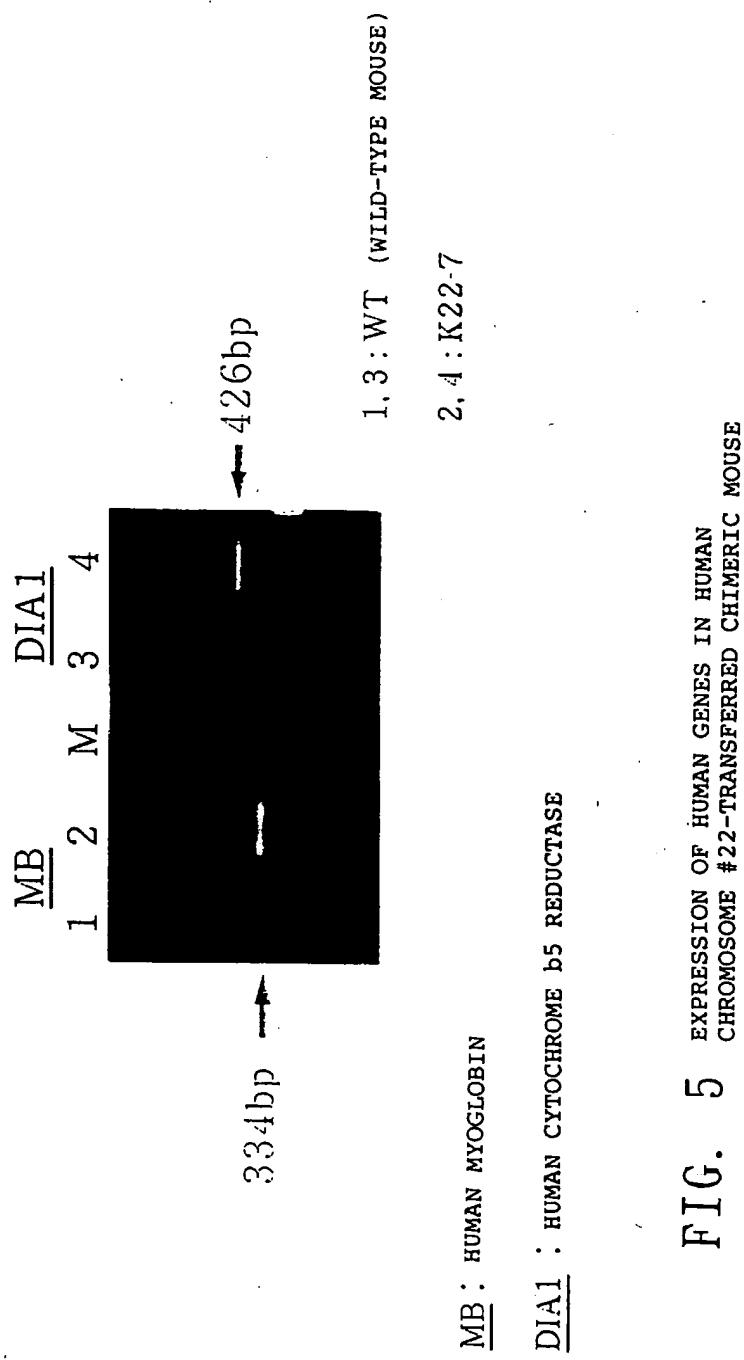


FIG. 5 EXPRESSION OF HUMAN GENES IN HUMAN CHROMOSOME #22-TRANSFERRED CHIMERIC MOUSE

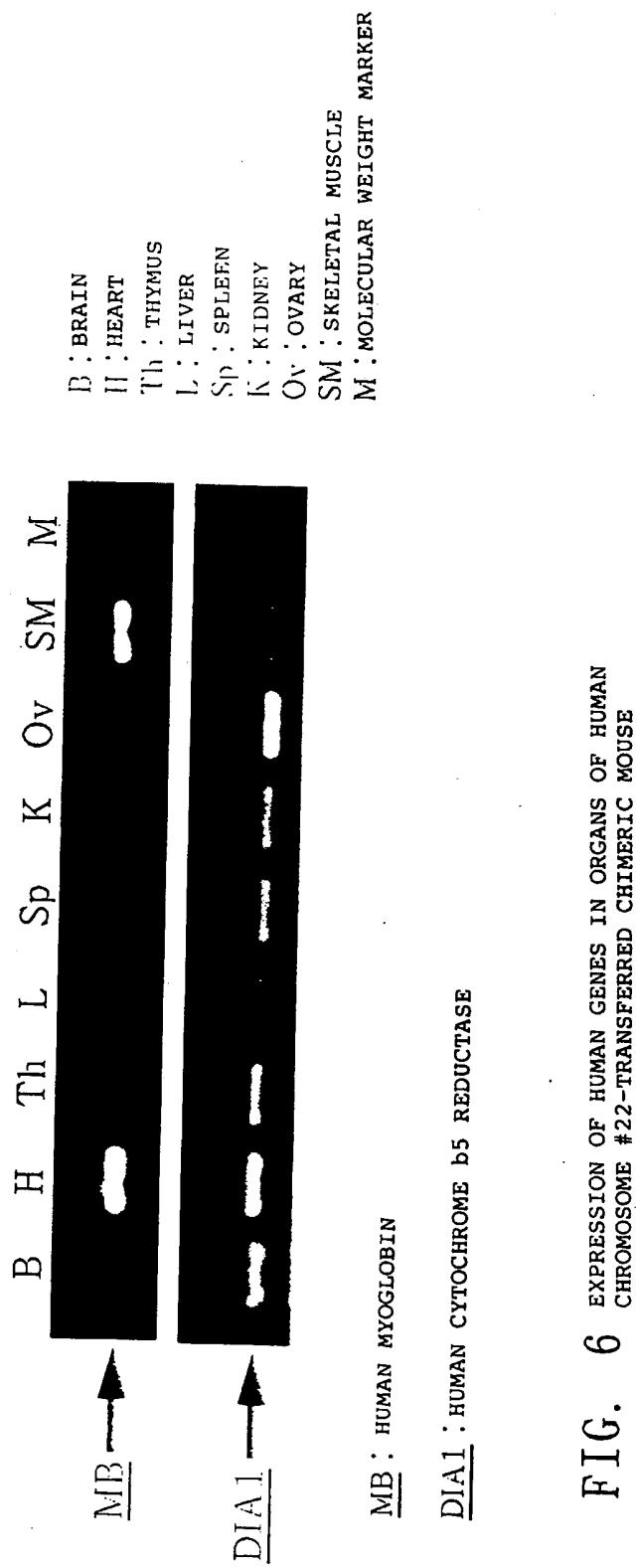
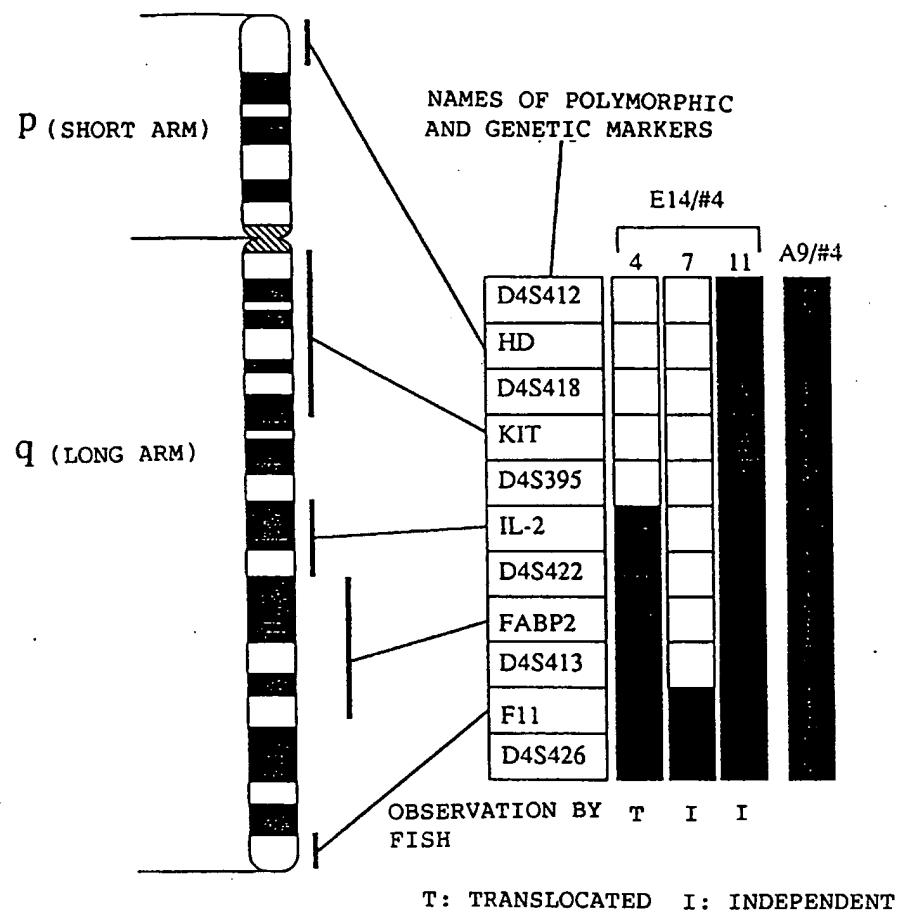


FIG. 6 EXPRESSION OF HUMAN GENES IN ORGANS OF HUMAN CHROMOSOME #22-TRANSFERRED CHIMERIC MOUSE



4

FIG. 7 RETENTION OF HUMAN CHROMOSOME #4 (FRAGMENTS)
IN E14 DRUG RESISTANT CLONES (PCR ANALYSIS)

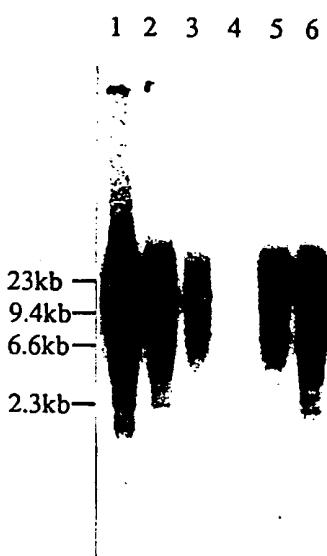


FIG. 8

DETECTION OF HUMAN LI SEQUENCE IN HUMAN
CHROMOSOME #4-TRANSFERRED E14 CELL CLONES (SOUTHERN ANALYSIS)

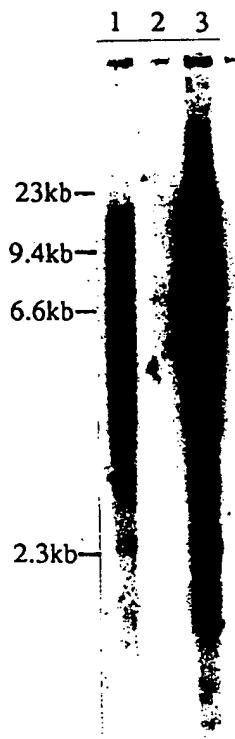


FIG. 9

RETENTION OF HUMAN L1 SEQUENCE IN CHIMERIC MICE DERIVED FROM
CHROMOSOME #4-TRANSFERRED ES CELLS (SOUTHERN ANALYSIS)

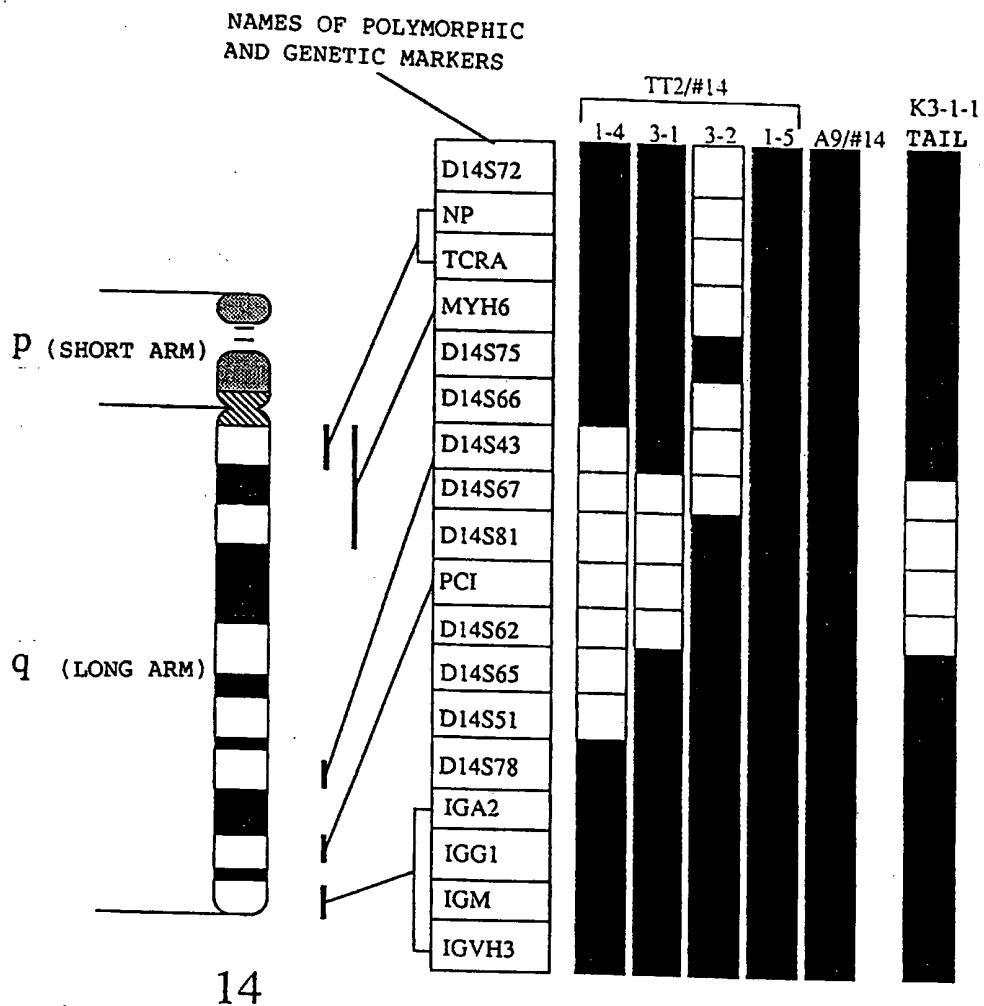


FIG. 10 RETENTION OF HUMAN CHROMOSOME #14 (FRAGMENTS)
IN TTS DRUG RESISTANT CLONES (PCR ANALYSIS)

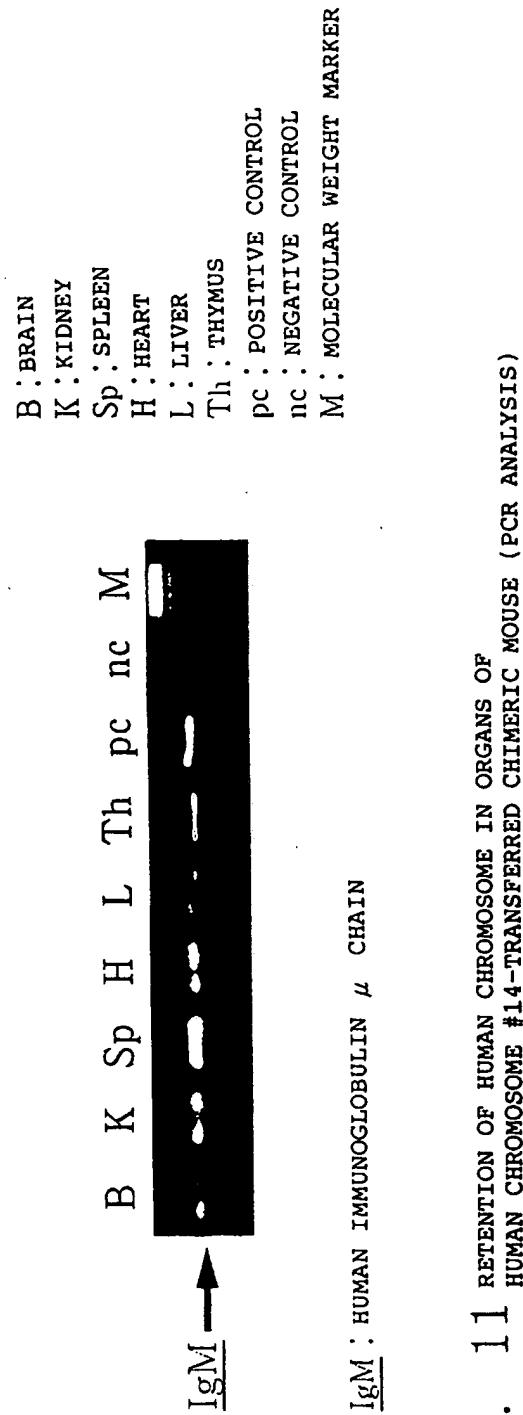


FIG. 11 RETENTION OF HUMAN CHROMOSOME IN ORGANS OF HUMAN CHROMOSOME #14-TRANSFERRED CHIMERIC MOUSE (PCR ANALYSIS)

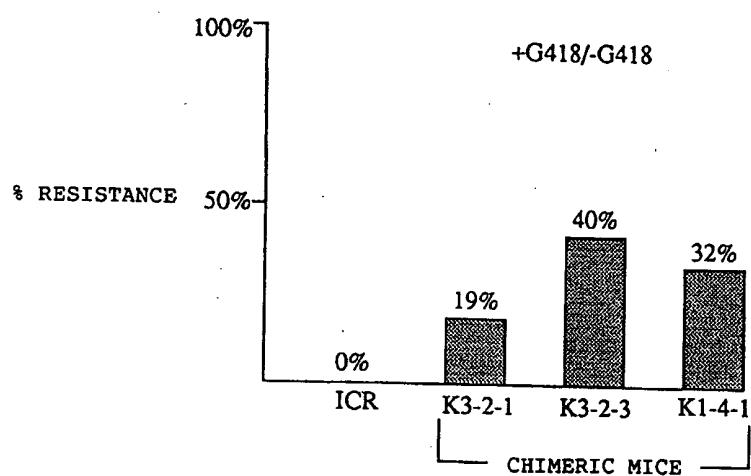


FIG. 12 TEST ON TAIL-DERIVED FIBROBLAST CELLS FOR G418 RESISTANCE

FIG. 13

HUMAN IgM CONCENTRATION IN CHIMERIC MICE IMMUNIZED
WITH HUMAN SERUM ALBUMIN (ELISA)

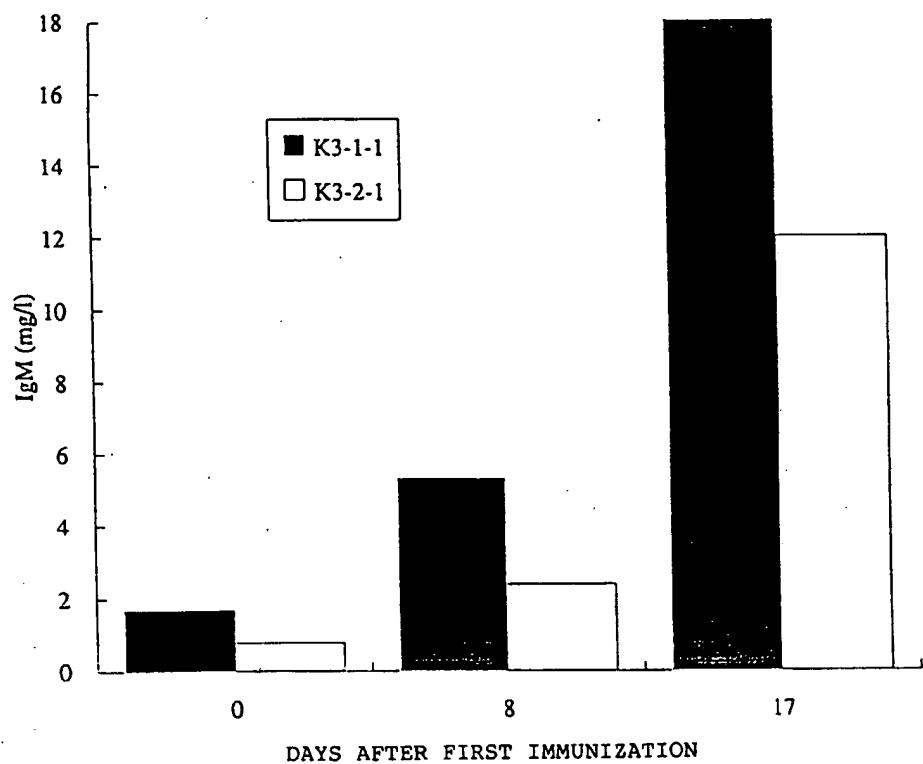


FIG. 14

HUMAN IgG CONCENTRATION IN CHIMERIC MICE IMMUNIZED
WITH HUMAN SERUM ALBUMIN (ELISA)

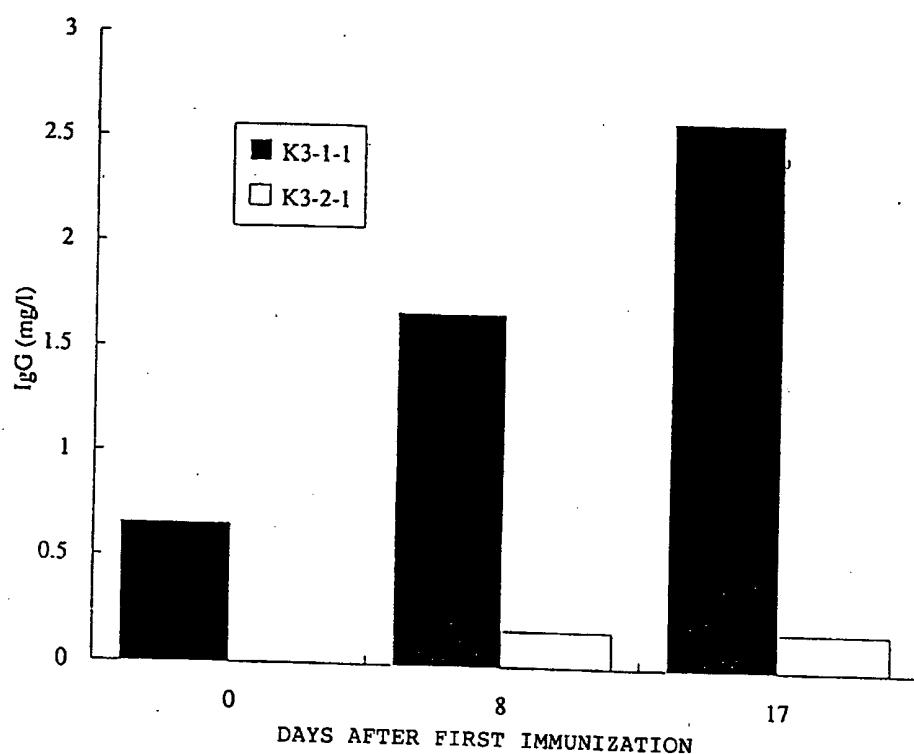
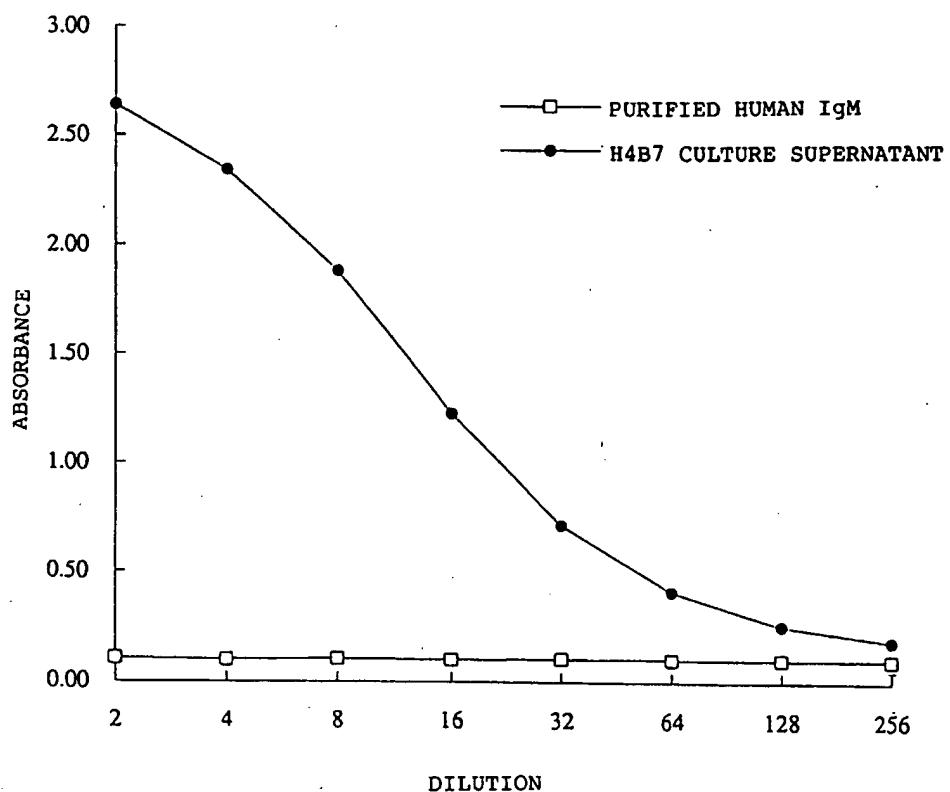


FIG. 15 ANALYSIS OF ANTI-HSA-HUMAN IgM PRODUCING HYBRIDOMA



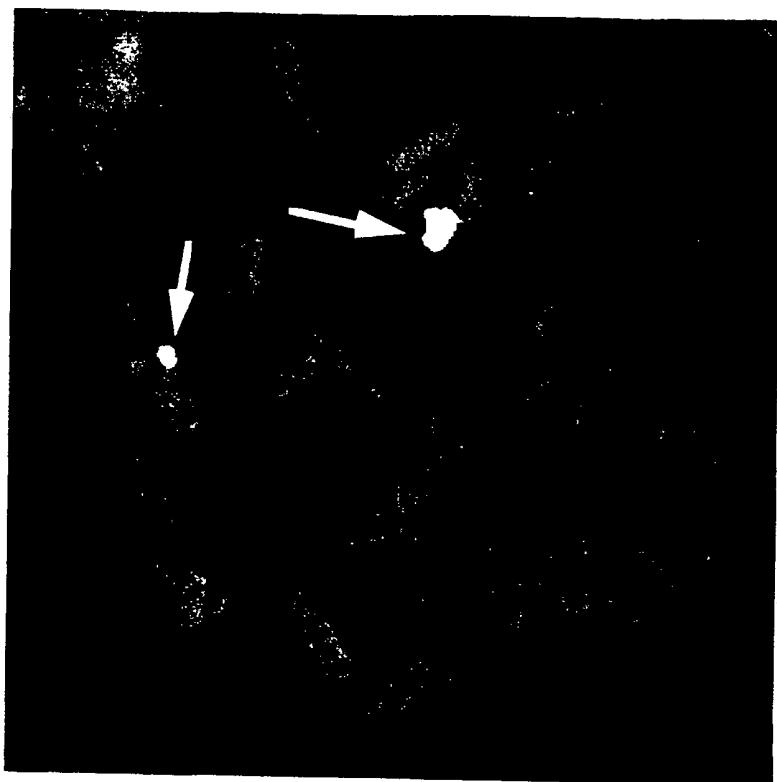


FIG. 16

FIG. 17 TITER OF ANTI-HSA HUMAN IgG IN SERUM
OF HSA-IMMUNIZED CHIMERIC MICE

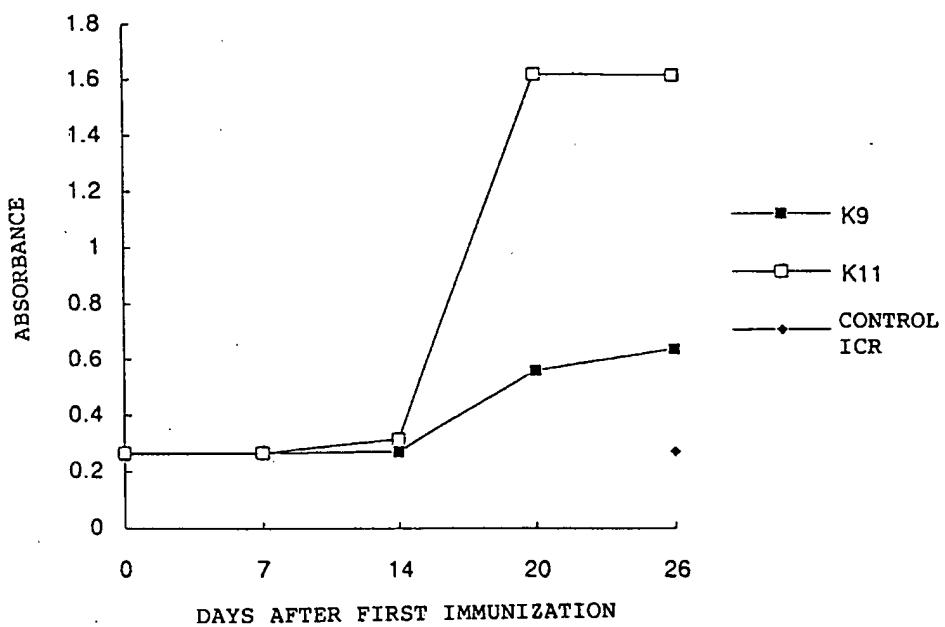


FIG. 18 TITER OF ANTI-HSA HUMAN Ig κ IN SERUM
OF HSA-IMMUNIZED CHIMERIC MICE

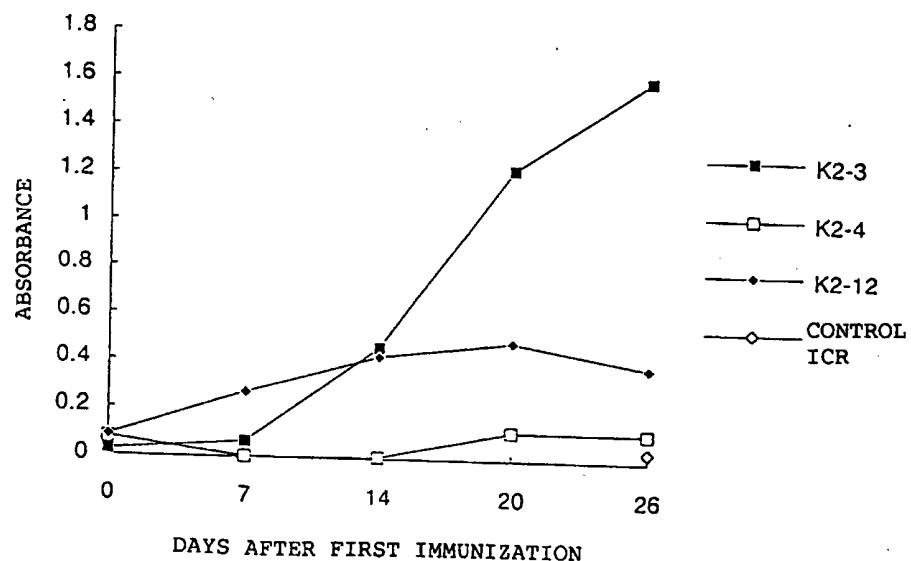


FIG. 19

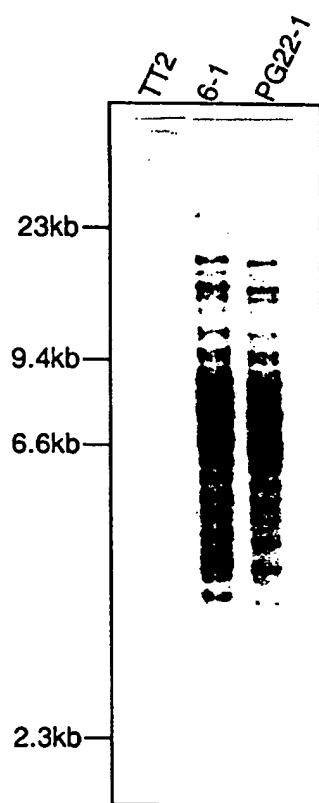


FIG. 20 TITER OF ANTI-HSA HUMAN Ig λ ANTIBODY IN A SERUM OF HSA-IMMUNIZED CHIMERIC MOUSE

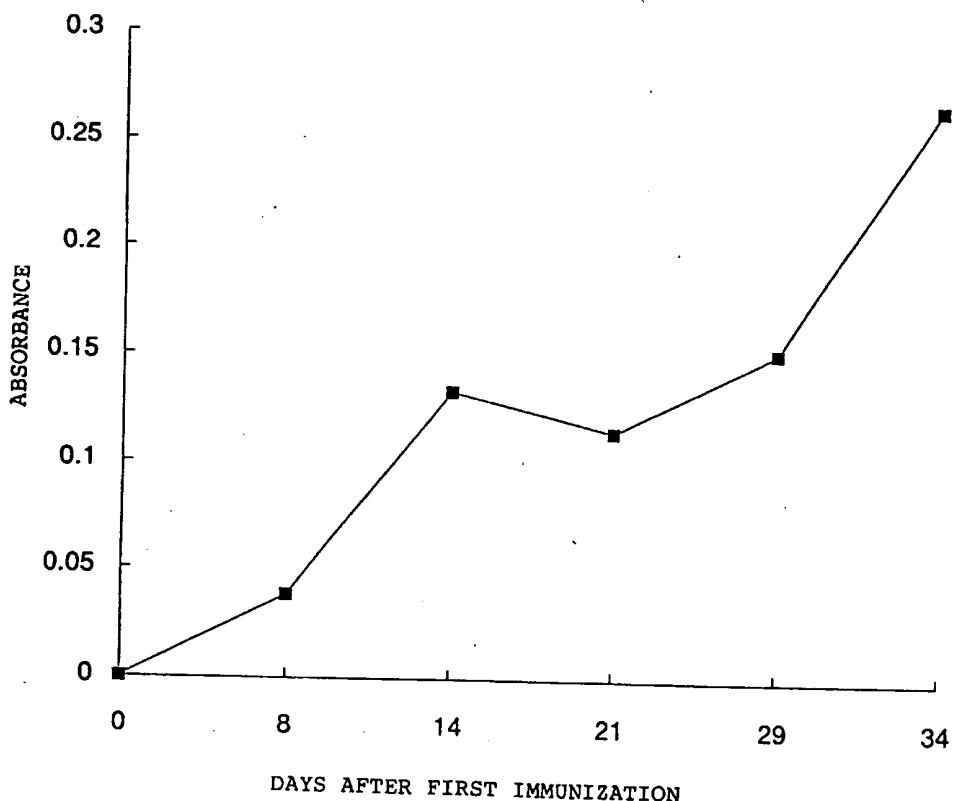
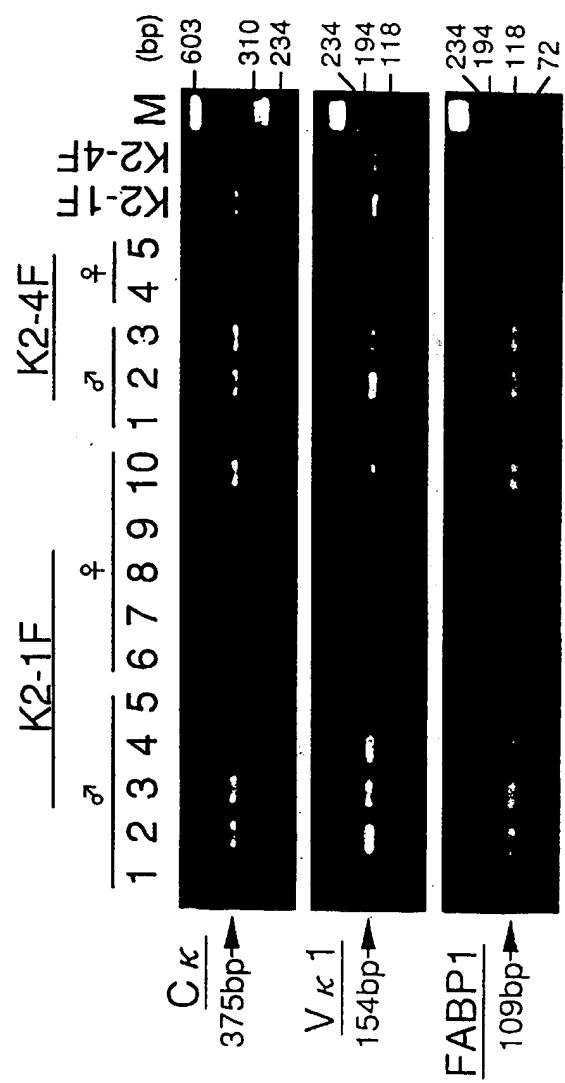


FIG. 21



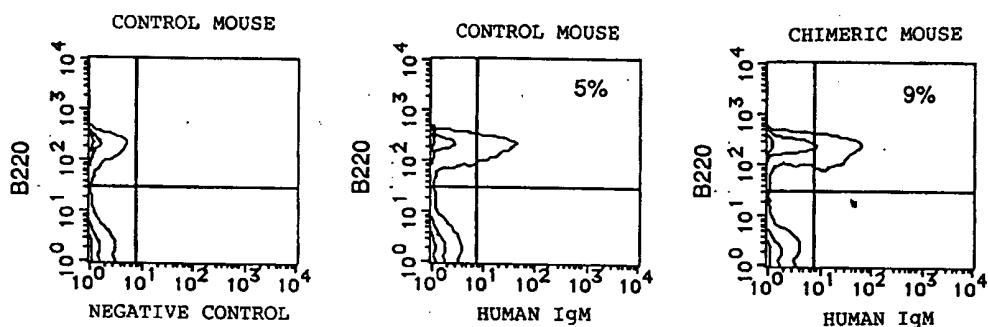


FIG. 22 ANALYSIS OF SPLEEN CELL FROM
HUMAN CHROMOSOME TRANSFERRED CHIMERIC MOUSE



Europäisches Patentamt

(19)

European Patent Office

Office européen des brevets



(11) EP 0 773 288 A3

(12)

EUROPEAN PATENT APPLICATION

(88) Date of publication A3:
09.07.1997 Bulletin 1997/28

(51) Int. Cl.⁶: C12N 15/00, A01K 67/027,
C12N 15/90, C12N 5/10,
C07K 16/00, C12N 15/08,
C12N 15/06

(43) Date of publication A2:
14.05.1997 Bulletin 1997/20

(21) Application number: 96113839.3

(22) Date of filing: 29.08.1996

(84) Designated Contracting States:
AT BE CH DE DK ES FI FR GB GR IE IT LI LU MC
NL PT SE

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(30) Priority: 29.08.1995 JP 242340/95
15.02.1996 JP 27940/96

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(54) Chimeric animal and method for producing the same

(57) Chimeric non-human animals retaining foreign chromosomes or fragments thereof and expressing the genes on the foreign chromosomes or fragments, and their progenies; a method for producing the chimeric non-human animals; tissues and cells derived from the chimeric non-human animals and their progenies; hybridomas prepared by fusion of said cells with myeloma cells; pluripotent cells containing foreign chromosomes or fragments thereof; a method for producing the pluripotent cells; use of the pluripotent cells for producing chimeric non-human animals; and a method for producing a biologically active substance, which comprises expressing the gene on a foreign chromosome or a fragment in the chimeric non-human animal or its progeny, or a tissue or a cell thereof, and recovering the biologically active substance as the expression product.

EP 0 773 288 A3



European Patent
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EUROPEAN SEARCH REPORT

Application Number
EP 96 11 3839

DOCUMENTS CONSIDERED TO BE RELEVANT									
Category	Citation of document with indication, where appropriate, of relevant passages	Relevant to claim	CLASSIFICATION OF THE APPLICATION (Int.Cl.6)						
D,A	NATURE, vol. 362, no. 6417, 18 March 1993, pages 255-258, XP002025628 JAKOBIVITS A ET AL: "GERM-LINE TRANSMISSION AND EXPRESSION OF A HUMAN-DERIVED YEAST ARTIFICIAL CHROMOSOME" * the whole document * ---	1	C12N15/00 A01K67/027 C12N15/90 C12N5/10 C07K16/00 C12N15/08 C12N15/06						
A	CYTOGENETICS AND CELL GENETICS, vol. 60, no. 1, 1992, pages 79-80, XP000671424 NING, Y. ET AL.: "Isolation of monochromosomal hybrids following fusion of human diploid fibroblast-derived microcells with mouse A9 cells" * the whole document *	1							
A	METHODS IN MOLECULAR GENETICS, vol. 1, no. part A., 1993, pages 115-133, XP000671521 LANGSTON, A. & FOURNIER, R.E.K.: "Preparation and properties of microcell hybrids" * the whole document *	1	TECHNICAL FIELDS SEARCHED (Int.Cl.6) A01K C12N C07K						
<p>The present search report has been drawn up for all claims</p> <table border="1"> <tr> <td>Place of search</td> <td>Date of completion of the search</td> <td>Examiner</td> </tr> <tr> <td>THE HAGUE</td> <td>16 May 1997</td> <td>Chambonnet, F</td> </tr> </table> <p>CATEGORY OF CITED DOCUMENTS</p> <p>X : particularly relevant if taken alone Y : particularly relevant if combined with another document of the same category A : technological background O : non-written disclosure P : intermediate document</p> <p>T : theory or principle underlying the invention E : earlier patent document, but published on, or after the filing date D : document cited in the application L : document cited for other reasons & : member of the same patent family, corresponding document</p>				Place of search	Date of completion of the search	Examiner	THE HAGUE	16 May 1997	Chambonnet, F
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THE HAGUE	16 May 1997	Chambonnet, F							